SEQ ID NO:	SEQ ID NO: of peptide	Mct hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13730	44098	Α	13810	80	952	RPGIVALNAILPLSRTRQSKPTR
		1				LKKRAIANCAPTWKKRFVALV
	ŀ					YRTA*RFPSITLSVAVT*PASAK
			1	i		NKAIKMTQKIEQSQRQERVAA
						WNRRAECDLAAFQNSPKQTYQ
		1				AEKARDRKLCANLEEAIRRSGL
						QDGMTVSFHHAFRGGDLTVN
						MVMDVIAKMGFKNLTLASSSL
						SDCHAPLVEHIRQGVVTRIYTS
						GLRGPLAEEISRAYCADSHHGP
	l .					FAMVNNMTDLTAQEPAWQTR
		1				DHLDDPVIGELRNRFGPDAFTV
	1					QATRTGVPVVWIKREQLLEVG
					1	DFLKKLPKPYVMLFDLHGMDE
						RLRTHREGLPAADFSVFYHLISI
						DRNRDIMLKVALAENDLHVPT
	1	1				FTKLFPNANWYERETWDLFGIT
						FDGHPNLRWIP
13731	44099	A	13811	92	735	RLLRPLPQPRSSVLPGLTTFHAS
		1	1			STSETRQSKPTRLKKRAIANCA
		1				PTWKKRFVALVYRTA*RFPSIT
		1				LSVAVT*PCVNIITIGCQVKTVT
		1				TIATAQIQRFTRFNHFSRFQYFR
		1				NSPKQTYQAEKARDRKLCANL
		1				EEAIRRSGLQDGMTVSFHHAFR
		1			1	GGDLTVNMVMDVIAKMGFKN
i	1	ĺ				LTLASSSLSDCHAPLVEHIRQG
		ı			l	VVTRIYTSGLRGPLAEEISRGLL
		1				AEPVQIHSHGGRVHLVQNGEL
		1				NIDVAFLGVPSCDEFGNANGYT
		ı			l	GKACCGSLGYAIVDADNAKQV
		١.	10010	ļ	1110	VMLTEELLPYTDNPEA
13732	44100	A	13812	5101	1113	LRRDSGRLRSCWHPRSPTSMGS
13733	44101	Α	13813	5121	6228	LGQREDLQDEDRNSGADPGQR
1		1				
		1				GRCSIHLEGPGQQPCLQRAVQG EGDPVLAKEEIQDQCHPHGQV
		1				OLLLVPAAEPVRAVPPRVOPVL
						PHHHHPAELSGIREE*GNQNDA
		1				EN*TISTIRTGSGLESSR*M/SILE
		1				LSRTROSKPTRLKKRAIANCAP
		1			l	TWKKRFVALVYRTA*RFPSITL
		1				
		1			1	SVAVT*PYWSGAPGGPCRRPYC HFRHRGARGSGAPGDGGEAPP
		1			I	AAAFLGGDVGFGL
13734	44102	A	13814	56	196	CRGDGGKLIIQCQRRMGI*PCG*
13/34	44102	A	13814	30	150	LRORKAELFOHLPLC*FWRPEI
		1	1		1	GTVCSGSIDNGSDRLRR
13735	44103	В	13815	64	2373	G I VCSGSIDNGSDRERR
13736	44103	A	13815	3	403	
13737	44104	B	13817	55	1040	
13/3/	J44103	I _D	113017	22	11040	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13738	44106	A	13818	587	1533	VILITAKFFFSMYSPIKTAGGFA STYVI-MISIDAEKAFDK HIQPF MLKALNKSENKIPRNPTYNGCI GPLQGELQTTAQGNKGYKQT GPLQGELQTTAQGNKGYKQT QQHDMHMSTKNQYHERNGHTA QGMKGITQPSAFIPTAESNSFOG QVKTLPSPIDAKQQLORKIQKK QQEQKLQSPLPGESAAKKSESA TSNGVTNLPNGNPSLSSPQFIGI VAAVPSPIPVQRTRQLVTSPSP MSSSDGKVLPLNVQSHQNCW WTRFNMSSLNVVKMTTISLTPS NSNTPLKHSASFISATGTTEESR SVPQIKNGSVVSLQSPGSRSSSA GGTSAVEVKVEPETSSDEHPVC QENSDEAKAPQTPSALLCQKS NTDGALQKPSNEGVIEIKATKE RKSSPATEGSWMENVFEGGRE GFRRSNHQIREDSOTIKKEEKE RKEELMELKTKVRELREECRSI RSEGODLEERVSAMDDEMNER KREGNFREKIKRNEQSLQEIW DYVKRPNLRLIGPPSDGEGNGL KLENTLQDIIQENFPNLARQAN QUQEQERTP
13739	44107	Α	13819	590	817	X-XX
13740	44108	A	13820	107	410	KGRNIQLNGPSSRVSLWRSLYF LNLNTGLPC*IGEVLLDNILQSV FQLGSILPITFRYTNQT*IWSFHI VPYFLEALLISFYSFFSKLPFSLF FIHFIFHR
13741	44109	В	13821	1	3735	
13742	44110	A	13822	922	1137	HFFLHFNFGESDNYVSWSFSSR GVSLWRSLYFLNLNVGLPC*IG EVLLDNILQSVFQLGSILHITFR YTNQT
13743	44111	В	13823	1	954	
13744 13745	44112	A	13824 13825	683	156 972	SHRLLRLLHSSRSSPGLVSSSISS FKHFSVLVILFTLMIVSFAVQKI FSLIRSHLSILALVAIAFGVLVM KSLPTPMSSMVLSR/SFF*GFYG FRSYI
13746	44114	С	13826	1	895	
13747	44115 44116	A	13827 13828	705	1727	LALLVELIPLPLCNGLLCLF*SL LV*SLFYQRLGLQPLPFFVFHLL GRSSSILLF*AYV*LFWHDFAA AGTGCSFPCLALPSGALVGQA WW*QKSLSICLSVKDFISPSLM KLSLAGYEILG

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13749	44117	Α	13829	1	1136	
13750	44118	A	13830	547	675	
13751	44119	В	13831	144	1877	
13752	44120	Α	13832	1	441	
13753	44121	Α	13833	27	344	
13754	44122	Α	13834	287	600	GPGSLQSSPLLSPPWPRASSPLL
		l				LRPSVRGWHLGEP*NCFPSPLA
		1				AS/EQAPKRSS*PQCNHR*RETA
			ŀ			ARGLAYPPDRPLPSMLSCESPE
						TPAKAGPGRERGCEKC
13755	44123	A	13835	2	226	
13756	44124	A	13836	2	226	
13757	44125	В	13837	45	518	
13758	44126	В	13838	1	699	
13759	44127	Α	13839	3	362	
13760	44128	Α	13840	2	130	
13761	44129	Α	13841	3	96	
13762	44130	Α	13842	1	545	
13763	44131	Α	13843	2	144	
13764	44132	Α	13844	2	91	
13765	44133	Α	13845	1	370	
13766	44134	С	13846	1	214	
13767	44135	Α	13847	215	310	
13768	44136	Α	13848	2	91	
13769	44137	Α	13849	1	289	
13770	44138	Α	13850	1	997	
13771	44139	Α	13851	1	192	
13772	44140	Α	13852	2	267	
13773	44141	Α	13853	2	181	
13774	44142	Α	13854	1	171	MESYAAIKNDEFMSFVGTWMK
	1					LETLILSKLSQGQK\TKHCMFSL
						TDVIARRRSWTSI
13775	44143	Α	13855	92	244	KRRRRRGQIRKMKEEEGNRRT
						WMKLETIILSKL*QRQKTKHRM
						FLLIGGN
13776	44144	Α	13856	146	382	
13777	44145	В	13857	124	1215	
13778	44146	В	13858	1	628	
13779	44147	Α	13859	377	484	
13780	44148	Α	13860	2	220	WARTSCLKHQKQWQQKPKLA
						NGI*LN*RASAQQKKLPSEWTG
						TWMKLETIILSKLLQGQKTKHR
						MFSLIGGN
13781	44149	A	13861	965	1150	GVDQRAVCSSRIW*SHLFLFHS
		ı		I		VVVHLPWAGTWMKLETIILSK
	1	1	1			LSQGQKTRHRMFSLIGGN
13782	44150	Α	13862	254	571	YSYKLEIIIGKQPEIYRDPCYRP
1	1	1	1	1		AV*SQTAVLAISEAPWA*DLPS
	1	1	1	1		QVSDTISWWTVC*AHWKSAVL
			1	1		GWE*PGFPGTWMKLETVSLSK
	1	1	1	I		LSQGQKTKHHMFSLIGGN

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	
1	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nuclcotide insertion)
				sequence		
13783	44151	A	13863	62	214	
13784	44152	Α	13864	274	435	
13785	44153	Α	13865	I	294	ASTAGVSYYVAQAGLKLLGLS
						LSKCRDYRCEPPCPE*MSLYKV
		l				MAMARKAMSL\YIYFFLDEFM
	ŀ	l		l		YFAGTWMKLETIILSKLSQGQK
						TKHRMFSLVGGN
13786	44154	Α	13866	663	1266	PPLTCQCSHLCVTLVGH/WIKLE
		l				TIILSKLSOGOKTKHRVFSLIGG
				1		AGALAKECMSLYPGCKITVFDI
		l		ŀ		PEVVWTAKQHFSFQEEEQIDFQ
l	l	l				EGDFFKDPLPEADLYILARVLH
ĺ	i	l		1	1	DWADGKCSHLLERIYHTCKPG
						GGILVIESLLDEDRRGPLLTQLY
		l	Ì			SLNMLVQTEGQERTPTHYHML
				1		LSSAGFRDFQFKKTGAIYDAIL
13787	44155	A	13867	1	681	(
13788	44156	A	13868	I	247	
13789	44157	Α	13869	2	178	
13790	44158	В	13870	1	2042	
13791	44159	Α	13871	1	378	
13792	44160	Α	13872	1	683	
13793	44161	Α	13873	ī	972	
13794	44162	Α	13874	308	406	
13795	44163	С	13875	177	653	
13796	44164	Α	13876	1	465	
13797	44165	Α	13877	3	1175	FFSSQAAALLVVLGSSHTLQRA
						RKHSAPGLPDTCALLQPPAASA
		l				AAAPSMSGPDVETPSAIQICRIM
						RPDDANVAGNVHGGTILKMIE
		1	İ		[EAGAIISTRHCNSQNGERCVAA
		l				LARVERTDFLSPMCIGEVAHVS
			ŀ			AEITYTSKHSVEVQVNVMSENI
		l				LTGAKKLTNKATLWYVPLSLK
						NVDKVLEVPPVVYSRQEQEEE
		l				GRKRYEAQKLERMETKWRNG
		l				DIVQPVLNPEPNTVSYSQSSLIH
		l	Ì			LVGPSDCTLHGFVHGGVTMKL
		l	Ì			MDEVAGIVAARHCKTNIVTAS
		l				VDAINFHDKIRKGCVITISGRMT
						FT\SN\NSMRFKVLLDPAPVVDS
		l				FQKRYRGRQCLSST*VSLSQER
		ĺ				QV/PLPVPQLVP\ETEDEKKRF\E
						EGKGRYLQMK\AKR\QGHAEPQ
13798	44166	Α	13878	121	36 I	SLFELENRSKFSVATAGLVPSR
						RLSINCIRRAFITSSNGFPLWFCL
		1				HQKGPVDCCVAVY*CGTEDGS
						WAPTHPGVGPSLP
13799	44167	Α	13879	1	462	
			<u> </u>			

A	13880	3	287	PCSLSSDCCASNQRDSVGIGPSE
A	i e			PRVGYNILVIRFLSPSEKRSPR\V GVTRFSSAWEPSQLAGILPISEP PSNRIFACWGKPAWTACCNSLR ARR
	13881	1	1332	
A	13882	1	753	
A	13883	1	516	
A	13884	71	277	IYFNTQESFCPPRPLK/LCDGRG CHKGL*NAFKAFSPFSWPLALG SFLLMQISAACLNSFPENGLFFS TT
В	13885	1	1095	
A	13886	1	741	
A	13887	264	2880	WILLOKHELHVSR*IGLIPDTSW ILLOKHELHVSR*IGLIPDTSW ILPSRPDAGGRFPIRTPPCVLVV HKPSRHCREWTANTVLPKGAL AKWLLEPKADLQGWHLPADG QVTKPSISYNIKLYINCYNRTVT TAAQHQQVQRKIIGASIFLAEN VAVREENBEERKYTRAPTEA WTAGSWIETAYREIKGVGQES VCITVMHLEELUYELLAWIIMSA VYEAFDPVAPSSKCELNPDPPD YCCTPYTLHTSPPPPPSCCTPYT LHTSPPPPPFSCCTPYTLDTSFS PQTHSCCTPYTLHTVPTFHLPA AHPHILHTIPPPGTCGTPYTLDTSFS PQTHSCCTPYTLHTVPTFHLPA AHPHILHTIPPPGTQGGFVSED VAAPDVRKVKITVNYPPYISKA KNTGVSVGQKGILSCEASAV MAEFQWFKEETRLATGLDGMR IENKGRMSTLTFFNYSEKDYGN MAEFQWFKEETRLATGLDGMR IENKGRMSTLTFFNYSEKDYGN MAEFQWFKEETRLATGLDGMR MFFETNKLGNTNASITLYVD HIVGSKALLSKCKRTEITNYLS KLNNPLLNDYWVHNEMKAEIK MFFETNKNKDTTYQNLWDAFK KLNNPLLNDYWVHNEMKAEIK MFFETNKNKDTTYQNLWDAFK KLNPLLNDYWVHNEMKAEIK MFFETNKNKDTTYQNLWDAFK KKREKPGIENKINGRILARLIK KKREKPGIENKINGRILARLIK KKREKPGIENKINDGLIARLIK KKREKPGIENKINDGLIARLIK KKREKNQIDAIKNDKGDITTYP TEIQTTIRESYYKHLYANKLENL EEMDKFLDTYTLPRLNQEEVSES
	B A	B 13885 A 13886	B 13885 I A 13886 I	B 13885 I 1095 A 13886 I 741

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
13808	44176	A	13888	1	629	AMAGSPPPASLLFCILISDCCAS NQRDSYGVOPSEPGVGYSLVV RRFLSPSEKRSIRVGVTRFSRCR PSPLSLTQKGNSLTSWASQVLT PAPEKQPDNLFLPRRGSRDPT EYFHGGEVRDKSDEHVDHVSQ HISFRDEVKRLGAEDSNAFUT TELHEFNTGGVYRPPNSTSVIQP LDQGVIRTFKPCYTKYSLEKIVS AVEENPDAT
13809	44177	A	13889	1	1452	
13810	44178	A	13890	167	309	
13811	44179	A	13891	3	386	SVGALQKQQGWHQRRCISLHF PSFLCVFSLAKSRRADHNQDPR ASRMAFSLTSSPLARSAASVA AFGCGTSGNGRFSSVP*AYRQV FQRIAIAVYGPLMRDR**AHPVK TPDAIDCRIRQPAIPRLQ
13812	44180	В	13892	121	298	•
13813	44181	Α	13893	473	1826	
13814	44182	Α	13894	1	270	
13815	44183	Α	13895	3	3706	
13816	44184	A	13896	1	1464	
13817	44185	В	13897	591	729	
13818	44186	Α	13898	1	1326	
13819	44187	A	13899	1	1152	

1705

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13820	44188	A	13900	1	2524	MREEFAHRWLARAGEKTPPDA
						LEKLTDPLMFAAAMVSAGKAD
						VCIAGNLSSTANVLRAGLRIIGL
		l			ŀ	OPGCKTLSSIFLMLPOYSGPAL
						GFADCSVVPQPTAAQLADIALA
						SAETWRAITGEEPRVAMLSFSS
						NGSARHPCVANVOOATEIVRE
						RAPKLVVDGELQFDAAFVPEV
	ŀ					AAOKAPASPLOGKANVMVFPS
		l				LEAGNIGYKIAORLGGYRAVGP
		l				LIOGLAAPMHDLSPCIAATDAG
						SAAAORIGDLVSVHVIPRPHVD
						LEOVFPIGLKAYTATCNHPVSL
						RFTIKSSSDRIKLSNQTLQSKNL
						FIMRFYCFSLCLSLSAFYHCRGV
				,		IVNIOAVFWLOKGTEAADILTT
						CAOLWIMMYALLSVFSLLDVIL
1						NLAOKFPAASOLPLKGIFOGIKL
İ	İ					IGAILVGILMISLLIGOSPAILISG
						LGAMAAVLMLVFKDPILGLVA
						VRNWDNTITTIPTWSLVSDSFK
						NWSGMSASGGRRIKRSISIDVTS
						IRFLDEDEMORLNKAHLLKPYL
1						TSRHOEINEWNROOGSTESVLN
						LRRMTNIGTFRAYLNEYLRNHP
				l		RIRKDMTLMVRQLAPGDNGLP
						LEIYAFTNTVVWLEYESIQADIF
1	i					DHIFAIVEEFGLRLHQSPTGNDI
						RSLAGVTINKAPGDDEILKNTV
						DFVSFSYYASRCASAEMNANN
						SSAANVVKSLRNPYLQVSDWG
						WGIDPLGLRITMNMMYDRYQK
13821	44189	Ā	13901	1	2370	
13822	44190	A	13902	1	282	
13823	44191	Α	13903	1	714	
13824	44192	A	13904	3	506	
13825	44193	A	13905	459	551	
13826	44194	Α	13906	401	492	
13827	44195	В	13907	479	702	
13828	44196	A	13908	120	256	

1706

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1	l		sequence		
13829	44197	A	13909	16	1249	SKSGAIEIARLAGRVHGVVVQI
				l		TSEIGPLTLLTANFAFSAS*STA
	i	l				WKATSMDGEVLS*YSTSASAS
	l	l				AERQSTHQ*TGFAPLCR*PLPMI
	i	l		1		/VAQRADDVGFSFEVHGQVRV
	l	l				RPVAOHAOTDKVFALTVNLGR
		l				RVFAALGAELGGARSAVRPVP
		l				
	l					ECYAVRETLALLQKIFPIRQCEN
			l			SVYRNRSRPCLQYQIGRCLGPC
1	i					VEGLVSEEEYAQQVEYVRLFLS
	l .	l			ŀ	GKDDQVLTQLISRMETASQNLE
						FEEAARIRDQIQAVRRVTEKQF
						VSNTGDDLDVIGVAFDAGMAC
	i .	İ				VHVLFIRQGKVLGSRSYFPKVP
					1	GGTELSEVVETFVGQFYLQGSQ
					ľ	MRTLPGEILLDFNLSDKTLLAD
	1			i		SLSELAGRQINVQTKPRGDRAR
		ŀ				YLKLARTYAATALTSNFRSIYR
				i	İ	SPATDRACQRVETAGNER
13830	44198	A	13910	1	438	3FATDRACQRVETAGRER
13831	44199	A	13911	i	1200	
13832	44200	A	13912	I .	300	
13833	44201	A	13913	1	431	
13834	44202	A	13914	2	1465	
13835	44203	A	13915	1	513	
13836	44204	A	13916	3	535	RDKILVFKDENFWMIRGYAVLP
13830	44204	A	13910	P	333	
1						DYPKSIHTLGFPGRVKKIDAAV
	l			i		CDKTTRKTYFFVGIWCWRFDE
		1				MTQTM\EKGFPQRVVKHFPGISI
						RVDAAFQYKGFFFFSRGSKQFE
				1		YDIKTKNITRIMRTNTWFQCKE
		1		1		PKNSSFGFDINKEKAHSGGIKIL
						YHKSLSLFIFGIVHLLKNTSIYQ
13837	44205	Α	13917	I	876	
13838	44206	Α	13918	23	414	
13839	44207	Α	13919	2	425	QERGPHQVIQRKECLREYSPVV
						S*AKGTGVAVGQKGTLQCEAS
						AVSSAEFHWYKDDKRLIEGNK
						GVKVENRPFLSKLIFFNVSEHD
						YGNYSCVASNKLGHTNASIML
						FGPGAVSEVSNGTSRRAGCVW
						LLPLLVLRLLLKF
13840	44208	A	13920	1025	1270	
13841	44209	A	13921	370	701	
13842	44210	В	13922	I	660	
	12.10	_	1.2.7	L ⁻		L

SEQ 1D NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nuclcotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13843	44211	Α	13923	1	432	RFSRVDDFVLTMDDGPPPRRAR
						GDLTHSGLWRVCCIEGIYKGHC
						FRINHFPEDNDYDHDSSEYLLRI
						VRASSVFPILSTILLLLGGLCIGA
			l			GRIYSRKNNIVLSAGILFVAAGL
						SNIVGIIVYISSNTGDPSDKRDE
						DK*NHYN
13844	44212	A	13924	1	1093	
13845	44213	Α	13925	1	1203	
13846	44214	A	13926	1	1353	
13847	44215	Α	13927	2	445	
13848	44216	A	13928	1	227	
13849	44217	Α	13929	1	840	
13850	44218	С	13930	120	356	
13851	44219	A	13931	40	342	
13852	44220	В	13932	403	473	
13853	44221	A	13933	70	2169	
13854	44222	A	13934	1428	1905	RCAATSGYGERVQRTV/PTHTV
		1			ł	W*ASLSQSSAHRVVAINSIANK
		1			ļ	RSSMPPKSWATVALTSTKLKCR
İ		1				FMAKVRNKSFSVLHNV*MTISI/
	ŀ	l		1		SPWVILSSGVDEKLFPRAVRVA
	l .			1	l	MTAGASGFLAGRAVWASVVG
		l		1		LPDNELMLRDVCAPKLQQLGDI
		L				VDEMMAKRR
13855	44223	A	13935	227	342	VHSWQRYSGR*RGY*PQNQSA
		ļ	10000	100	000	TDQTRRWQSLKTAGAVA
13856	44224	A	13936	480	928	FFADLQQPVCADSYPLLVQMK
i					İ	LAPHKS*NAVLWQRSATKSFSV
		l				L/AQGLNDHINMPWVILSSGVD
						EKLFPRAVRVAMTAGASGFLA GRAVWASVVGLPDNELMLRD
		1				VCAPKLOOLGDIVDEMMAKRR
1						FIPLLRRWVDLALTRWLITVSK
13857	44225	A	13937	564	1185	WLHASAALLRW*YOLPGOPHH
13857	44225	A	13937	364	1183	RRCRSADGSHVPSLCAAVPDQR
	l					\PLITVRRIGMQAMTTSTTGDRQ
						RSWOIAMDGSOKLPORMLDSV
						RWHLAHDSKFDLLALGVAGW
1	1	ŀ			1	MRYVGGVDEOGNPIEISDPLLN
		ŀ				RMAANRFVHITOLPVIYRAFRN
						SFTQVARLQLCHKRFQQRHTSV
	1	1			1	REQIVTVLSGIRDNHVHFVQTL
		İ	l		I	TGDGVGNQRQLVQ
13858	44226	A	13938	228	424	HRFCPAARH/RSWSGLREMAVS
13838	44220	A	13938	220	424	ISMPRRIINSGRYCAICWYKRRS
		l				RCARNMLIFPVARSTPIPGPS
13859	44227	В	13939	128	1670	RCARIVILIEF VARSTEIFGPS
13860	44227	A	13940	2	2367	
13,000	44228	lu-	13940	<u> </u>	2301	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13861	44229	Α	13941	ı	921	ELPASPAPCARTPOPLGGRWEA
						ARRKER VLCLFD VDGTLTPAR/
						QGRTLKIGSSHRECHADTPEAE
	ŀ		Ì			CGVCSSVTEVRAKWRSQTC*KF
1						ECYASPLPP*KIDPEVAAFLQKL
	1					RSSYLPVGSQTWSHTLSSSCPLC
			l			DPEVIGRDTVPVSGSGSPGHAQ
						SPVFLGQTIQNHLGPGGARFEF
	ŀ					RA/CSGSLWLPALPHH\RGTFIEF
						RNGNCVPSQGLAHSRSSELPSA
	ŀ	l	İ			PSPLQKEQAALLLALSAPPSRE*
l		l				QSPPLPTGGMISFDVFPEGWDK
						QVRSSQCGEGLHLNIPPLPILCV
		İ				QGGNDFEIFADPRTVGHS
13862	44230	Α	13942	105	416	HSVLEVGPWWEVFGSWGQILH
					1	SLVSSHHVRHWLLPSPSAMTES
	l	l				SLTFQQKPSRCQHHASCKAYHS
ł		l		1		HQLLLVPTAGRLEDRSHH*IPC
				1		KHSPAPAWSLAAPVGN
13863	44231	Α	13943	191	573	SSSGGYQQQLMRVIGTRDTAV
l						NKASNVPSFGA*I/PSGRGQQAK
l		i				LIALTRALTLAKELHVNIYADC
		1				KYAFHILHQHDVIWAERGFLTV
						QGSSSIINASLIKTLLKATLLPKE
						AGVIHYKGHQKASAPIA
13864	44232	Α	13944	1	1065	
13865	44233	A	13945	321	537	
13866	44234	A	13946	346	1260	GGPARAEFSRHLGCKGHHPLPK
		1				DVASEVPAGSSTSVSSLIRTNGC
ļ		1				PGLGSWEPP*FFSENKRLSTLMS
		ı				FTLRDSKMFSPQSGLFLGCSAG
l		1		İ		EHSALRDPTASGLPGTASGVRP
		l				GR/RQRVGSDIEASTQGESKPCI
		ı				LSI*TLPNA*LKTKTLTPLEIAAC
				1		TNPHSFSITARIGPAHLAAPRPV
l		1				TPAGRGGARAPGR*CAAARPPP
l		1				ASQ*VSGSPAPGGGPSYAPAAR
						TS*PSSLAPGAKRLRFSRVATFC
	l		l	l		CSFLAWSSSSNKAFCT*EHMSL
1				ŀ		LSSWISQSLEKVASCCPCEQFDS
L		L.	L			WKRLENILLFS
13867	44235	В	13947	1	459	L

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide		Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ļ	sequence		
13868	44236	A	13948	63	1263	VEAAKKAHHAACKEEKLAISR
						EANSKADPSLRPQAHTYALSPL
						FLTAMPSV*QGEHVIYTRVRHS
						DQVSESSGKPLSLPLLEPGIWAF
						EAAQVPPRWDQVRSLEPTTEW
						EAPGLKKRRVSQGLNGNEPLSC
						AWIFL*TKEKYEKSLKELDQGT
						PQYMENMEQVFEQCQQFEEKR
						LGWDKG*LAPAF*KSHVSL*FS
ł						YKAIYHDLEQSIRAADAVE/DR
1						RGF\RK**TFGFSPSAPVA*EWS
						ADLNRTLSRREKKKATDCQDP
ŀ						HGPLQSWDPLKALSLHSTLNVP
						SNPAQSAQSQSSYNPRGPHTLC
						RPVLRALWPPLSCSVSSYEKTQ
		İ				SYPTDWSDDESNNPFSSTDANG
1				1		DSNPFDDDATSGTEVLCSPPSV
	Ì			1		SYLSWRRLPLCLAGDELTKME
						DEDEQGWCK
13869	44237	Α	13949	2	485	AAAFGTRLSRRGRGCGVTGSG
ļ						GGRRRVPAPRRPRSPLQPPHQ/P
1						PRRPSRRTPGCGSRGA\PSRGRR
						LVPPPPEPPRAAQQRARH\HRTS
1						\RPAARQPPRRPPAQPRSCPRRV
						RPTPAWSPPGRRGPRRSRRRR
						RGGRARARRPRRRGQVPRPQS
	L	_				ATGAGHAR
13870	44238	A B	13950	1	207 5297	
13871	44239	В	13951	1206	1360	
13872	44240	A	13952	46	369	NEGKNAKGSORERSGYPOREA
138/3	44241	^	13933	46	309	HOTNSGSLGRNPTSOKRVGANI
l		1				OHS*RKEFSTOHFISSOTKLHK*
						RRNKILYRQANAERFCHHQEGS
-	l			ı		TKHGKEQPVPAAAKSCQNVK
13874	44242	A	13954	476	868	KYDCCISSTFCRALVHFSRTTFO
13874	77272	l^	13934	[""	000	VNSPLSPNISAFGYMPLKYSIYF
						LHHLRPNIL*AASYFLKYKGTL
						KLPALRGCLS*STRCWCKIHICI
1						*ND/ECLQFLGPWFENEFCFLPS
1	1					TSOFVEFVLDILVLONEP
			L	L	L	13Q1 TEL TEDIE TEQUEF

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	acteurs, repossible nucleonae insertion)
13875	44243	Α	13955	1770	3003	PANIIMTGS\NSHITILTLNVNG\
		1				LNAPI*RHRPAIA/S/WLKSQDPS
				1	İ	VV\CiQETHLTCRDTHRLEIKG
		l				WRNIYQANGKQKKAGVAILVS
İ		I	1			DKTDFKPTKIKRDKEGHYIMAK
		l				GSIQEEELTILNIYAPNTGAPRFI
		l				KQVLSDLQRDLDAHTIIMGDFN
1			l		1	TPLSTLDRSTRQKVNKDIQELD
		1				SALHQADLIDIYRTLHPKSTEYT
		1	ŀ			FFSAPHHTYSKTDHIVGSKALL
		1				SKCKRTETITNCLSDHSAIKLEL
		1				RIKKLTQNHSATWKLNSLLLND
		l				YWVHNKMKAEIKMFFETNENK
						DTAYQNLWDTFKAVCRGKFIA
		l		1		LNAHKRKQERSKIDILTSQLKE
1		1			1	LKKQEQTNSKATRRQEITKIRA
1		l		ł		ELKEIETQKTLQKKQPNKDPAM
		l				EKNNKTDRVLARLIKKKREKN
						EFSAHILYKFSKQNGS_
13876	44244	Α	13956	105	392	RSRKPRGNPKSHPEDTDPTKAP
		l		ł		RKRQISSRGHRSHR*EPGPQSAA
		l		İ		PGFREQRRRGQMG*IWYLTPTP
1		ı		İ		PGGQRPSSAASPRGFPPTNNSSR
		_				LPAEPE
13877	44245	Α	13957	55	403	SEGMSGKASEKQEAFACPRHLT
1		1	İ			RAVDGPS*SGLLGSRLPPSCSPG
		1	İ			KEAPGGSELYLCLMRGLQTL*A
İ		1				RSGTGLR/PNPAEIR*EPGPQSA
		1				APGFREQRRRGQMG*IWYLTPT
10000	11016	ļ	12050			PPGGQ
13878	44246	A	13958	217	444	RSRKPRGNPKSHPEDTDPTKAP
		1	i			RKRQISSRGHQSHR*EPGPQSA
				1	i	APGFREQRRRGQMG*I*TPCRP
12000	11017	١.	12050	309	670	ALPKSKPRPL PCPRASSQGLKATH\GQR*SSEP
13879	44247	Α	13959	309	0,0	PGASFPGLQDGGRREPRRPD*D
						GPSTGLVRCLGHANVGDPCLK
1	1	1	1	l	I	
		1		1	1	VFVFFKNLSPTYLRASAEQTLP
				l		LLLPHLHGLCLHQPLHLGFTAA
12000	14248		12060	202	760	WAQLTFWEASQ
13880	44248	С	13960	392	760	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13881	44249	Α	13961	24	915	HAGRHARGSLFPGARPSETRAH
		1				CVRTARPNCKEEDTPMVRGKN
		l				SGGEHCTWGPSLAAGG*GIGDP
		l				GGLPGT\PHPGLQLLLHGFAGP
l			1			HMASLCCGVSSPAGVRQWPA*
			l		1	GSGNHQVKSWLGLPKNVS*AP
					İ	SML*NPKMQRLMKAQPHGRCE
		ĺ	l			VRRGRVCSALGP/IGMSGKASE
						KQEA*DEGHPHLHAPGT*QDQ
1						VDGPS*SGLLGSRLPPSCSPGKE
		l				APGGSELYLCLMRGLQTL*ARS
						GTGLRANPAEIR*EPGPQSAAP
						GFREQRRRGQMG*IWYLTPTPP
					_	GGQRPSSAASPRGFPLV
13882	44250	A	13962	397	612	RPQHHTFCCDGRVMMARQKGI
						FYLTLFLILGTCTLFFAFE/CSFG
		Į.	ļ			*VELPDYGYFGSRGRTCFLGSL
						GLVLG
13883	44251	A	13963	125	1298	FQSWAQPLFLLSCNRKTHFGTR
						IPI\MSVMVVRKKVTRKWEKLP
			l			GRNTFCCDGRGMMARQKGIF\
		ĺ				YLTLFLILGTCTLFFAFE/CGRYL
						AVQLSPAIPVFAAMLFLFSM\AT
						L\LRTSFSDPGVIPRALPDEAAFI
		l				EMEIEATNGAVPQGQRPPPRIK
			l			NFQINNQIVKLKYWYTCKIFRA
						SRASHCSICDNCVERFDHHCPW
						VGNCVGKRNYRYFYLFILSLSL
			ŀ			LTIYVFAFNIVYVALKSLKIGFL
						ETLKETPGTVLEVLICFFTLWSV
			l			VGLTGFHTFLVALNQTTNEDIK
					ł	GSWTGKNRVQNPYSHGNIVKN
						CCEVLCGPLPPSVLDRRGILPLE
						ESGSRPPSTQETSSSLLPQSPAPT
						EHLNSNEMPEDSSTPEEMPPPEP
13884	44252	A	13964	44	647	PEPPQEAAEAEK DYSAQHGKASQKHVARLISSGT
13884	44252	l ^A	13964	44	047	
						FQGATVSRPFQLSTSSSSFCSSSS SLSSSSSLSSSLLLSCSSESVFLS
İ						LSFFFCSEASFLPFCSRLYASRA
		l				
						SFNGVTNRWNSISSMAESTSLA
	1					FSVFRFPFIAKLLAQDVAYSTKT
				1		LAAREIGAPGLLGDVDTVRELR
-	l					GDRG*AKLPFQHLLPSCPFVRPP
12005	44052	<u>_</u>	12065	160	1076	RAAHRLLDDPGDHGIGQV
13885	44253	В	13965	158	1075	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			l	sequence		
		<u> </u>				
13886	44254	A	13966	125	1297	FQSWAQPLFLLSCNRKTHFGTR
		1				IPI\MSVMVVRKKVTRKWEKLP
		1	l			GRNTFCCDGRGMMARQKGIF\
1	1	1	l			YLTLFLILGTCTLFFAFE/CGRYL
1	1	1	l	1	[AVQLSPAIPVFAAMLFLFSM\AT
			l	1		L\LRTSFSDPGVIPRALPDEAAFI
1		1	l			EMEIEATNGAVPQGQRPPPRIK
1		ı	l			NFQINNQIVKLKYWYTCKIFRA
1		1	l			SRASHCSICDNCVERFDHHCPW
1		1	l	1	[VGNCVGKRNYRYFYLFILSLSL
1			l	1		LTIYVFAFNIVYVALKSLKIGFL
1			l		ļ	ETLKETPGTVLEVLICFFTLWSV
1	1	1	l	1		VGLTGFHTFLVALNQTTNEDIK
1	l	1	l	1		GSWTGKNRVQNPYSHGNIVKN
1		1	l	1		CCEVLCGPI.PPSVI.DRRGILPLE
1	1	1	l	1	[ESGSRPPSTQETSSSLLPQSPAPT
1		1	l	1		EHLNSNEMPEDSSTPEEMPPPEP
1		1	l	1		PEPPQEAAEAEK
13887	44255	Α	13967	247	850	DYSAQHGKASQKHVARLISSGT
1		1	l	1		FQGATVSRPFQLSTSSSSFCSSSS
		1	l	1	j	SLSSSSSLSSSLLLSCSSESVFLS
1		1	l	1]	LSFFFCSEASFLPFCSRLYASRA
1		1	l	1		SFNGVTNRWNSISSMAESTSLA
1		1	1	1		FSVFRFPFIAKLLAQDVAYSTKT
1		1	l	1]	LAAREIGAPGLLGDVDTVRELR
1		1	İ	I]	GDRG*AKLPFQHLLPSCPFVRPP
			l	1	ļ	RAAHRLLDDPGDHGIGQV
13888	44256	В	13968	298	380	
13889	44257	A	13969	48	252	QRAGSPHSPRSLSAPPLPELPLW
1		1	1		1	RHLRSPSAHRCTVGAPFWDGQ
1		1	l		l i	G/LEPAPSAALAAFPRSRARDLQ
1				1		LA
13890	44258	В	13970	1	447	
13891	44259	Ā	13971	236	559	MWLEPMQMGFLHMMEKMAA
1		1	l	1		RTSAILD*GTLK*FHFTLTTSLK
1			1	1		ALSSHTPIFPGTGELQLPVSPSV
1		1		1		CLDQGMQLKPSTSSHLLKTVKP
1			1			RMKRQSLLHMKQSFEPKIYL
13892	44260	A	13972	1	645	MKPOTLTVSVTAFKVAHLEFVP
1	1	ľ	l	1		SDVQMCSEFLPSGGFVVSLASG
1			l	1		VKLOTFAVSFTAHKSSVDPKNS
1			l	1		EAQLASPSGFCTRAADGAACQS
1			l	1		LPCTCTPOPLGGRWDWALWS
1		1	l	1		RGW/LLVEEA WAAQEPTEELRE
1		1	1	I		RALALASPERGSHSAAVGORAP
1			1	I		QVPPKWEPRQRRRRERARAVR
1		1	1	I		TASTLSPLNPPSKQDTLTAVGN
1			l	I		LADDHSSYFLLDRGEEGALQL
13893	44261	A	13973	28	322	DADDII33 II DEDRUGECUALQE
13893	44261	A	13973	1760	4426	
13894	44262	B	13974	942	3414	
13033	144203	$_{\rm p}$	13973	744	2717	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13896	44264	Α	13976	1	274	
13897	44265	A	13977	545	1150	RISGPGAVLLPVMPALWRLRHR MFSLTKK*HLPAHRSLAEGYLL VPQPQLDTLKARLTJGKYGELQV LFEAYQIKKTKLDRQSSSASLET LLALLQAEGAKIEEDTEMMAEK FLDGELPLDSFIDVYQSKRKLA HMRRVKIEKLQEMYLKGQRLXS TRPWPRCPDGCPNWHLPPPFFT LPQSQWASCRCTSAHPPPTTPG AAGRIL
13898	44266	В	13978	213	611	THORE .
13899	44267	A	13979	1	639	MEAVTVVPVGTEEGMGEEEQD SGTTTGCGLPSVEMLATNPOK TIPISLLQKYGTRIGK\TPVYYLL KAEG\HQPNFTFW\V\GNTSCT AQQPRKKVVKHKAAEVALRHL KVESMLEPAPDESSFSPL DSSL FEDFPVFTAAAAATPVSSVFLT RSTPMEKQPPLLP\QSEC\PVG ALQKLVVQKGW\PLFHTVT\B SRPDQHEELT\RVERFT
13900	44268	A	13980	361	894	AFFPMSTAGVCGNVEGEPETP WSLPLSQVPFSPNA/CPQGTLGR S*AGVAGAPGPGVPGRSPAA/GI PEPAARPGRPALSRLPGLGGAG ARSGAA*G*GSRPGG/PLPAADP VQSWGP*EAPSHP*PGT\SADSP PSGVSGATGDCLWGSGRAELF GFATTSSAAPFPGLPTFFIPVSP
13901	44269	В	13981	1	3842	
13902	44270	A	13982	3	407	DAWAAAROGRSCALPPPGA/PE EPGHVPGAA*G*GSRPGGRFQL PDPVQSWGP*EAPSHP*PRNFPQ TRRHLGVKQHGCSQPSPHGSRA SQAHPGISGLF/GPHHPGPPLFIH GPRCCCAQATASNHLFACGTSS DP
13903	44271	С	13983	127	297	
13904	44272	Α	13984	3	1920	

1714

AGAMEAEROPERPAERSS TPEEGAQALAEFAALHVWS GIGVGSPAHTPADONSCPGV AGEVFGIMQVEEVEEEDE, EVRKQOPPHEAPTSLCPM QPHIVACARSIFLIDHAWTC EHARQQLQVPGLHRMAI MGIEHGELPINVPEGSSGG MGEREVPYVLCLQTAEEKN WYIMDEFGSSIG(HADVPSF) PUEEGPQPSVSPPHLMGTH/ GTEEVTRDFAYGETDPLIRK LPLSSSHRGMETTCVHTSRA PLQACELPSWSLLLINEPPRE PSLP*HLYTRCVSF*AIPGLI SRLDTWEPGSPSGGSPHRVY VQVASSLTHPALPVRASRI PPLELAHLPPCRKLSQERFW LINGPFCENLLTVKDCLASIA AGGPEGPPWVLGWHILG, APGAPQTALLLPRGEDNIW PWNLGLCVLGSPRGLCPGH ACTCVLQVVSKYESPVLEL DVGKVKFDIRYIVGGTSLAC PLKNPYVSAHLSSRAFAQG RGGPTWSGWLLARDPARV- LSSLVHSVERCPMQWALSE LHHLSTWSSGLGGLAG**VF HRTTPTP**QAACSRPSGGL A*WALCLLPGAADPLLRGC AS**LRLBGPQAEIFRAFTELF KTGIQKYRREALLQHYLGP, AGRSKALTGTAERSCSSGSE FTLCVNYNDSRAIMKTLAAI GFDCASKAEIQLVQILGMPP FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIIS PPCHISVKSGAMLKASRLLJI KEL 13907 44275 B 13987 1 2040 LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C LYFLETGVFCVQVPAQQC**C	SEQ ID			SEQ ID NO:			Amino acid sequence (X-Unknown,
13905 44273 A 13985 3 2271 GQGRRTAGGGGGGAGVG AGAMEARGPERRPAERSS TPEEGAQALAEFAALHVWS GIGVGSPAHTPADGNSGFQ) AGEVFGIMVGEVEEEDE, EVRKQQPNPLAPFISLCPMF QPHIVACARSIFLIDHAWTC EHARQQLQVPGLLHRMAI MGIEFHGELPIIVVPEGSSGG MGREVEPVLCLQTAEEM, WYIMDEFGSRIQHADVPSF, VPLEEGPQPSVSPPLMGTH, GTEVYRDFAYGETDPLIRK LPLSSHRGMETPCVHTSRA PLQACELPSWSLLLINEPPRF PSLP*PHLYFRCVSF* AIPGLI SRLDTWEFGSSGSPHLYVVQQVASSLTHFALPVRASRI PPLELAHLIPPCRELSGERF LNQFPCENLLTVKDCLASIA AGGPEGPWWLGWHHLG APGAPQTALLIPRGEDNINH PWMLGLCVLGSPRGLCPGH ACTCVLQVVSKYESPVLEL DVGKVKFDIRYIVGGTSLAG PLKNPYVSAHLSRAFAQGI RGGPTWSGWLLARDPARV* LSSLVHSVERCPMQWALSE LHHLSTWSSGLGCLAG**WY HRTPTPP**QAACSRPSGGL**QA**WALCLLPGAADPLLRGC AS**LRJPGPAEFFF**LFT**CFT**CFT**CFT**CFT**CFT**CFT**	NO:		hod				
13905 44273 A 13985 3 2271 GQGRRTAGGGGGGAGVG AGAMEAEROPERRPAERSS TPEEGAQALAEFAALHVWS GIGVGSPAHTPADGNSGPQ AGEVFGIMQVEEVEEEDE, EVRKQOPNPLAPTDSLCPMP QPHIVACARSIFLIDHAWTCI EHARQQLQVYEGLLHIMAT MGIEHGELPINVPEGSSGG MGEREVPYVLCLGTAEEKN WYIMDEFGSRIGHADVPSF, VPLEEGPQPSVPSPPLMGTH/// GTEVTRDFAYGETDPLIKK LPLSSHRGMETTPCVHTSRA PLQACELPSWSLLLINEPPRE PSLP*PHLYFRCVSF-AIPGLI SRLDTWEPGSPSSGSPHRVY VQQVASSLTHPALPVRASRI PPLEAHLPPCRKLSQERPE LAQFECENLLTVKDCLASIA AGGFEGPBWVIGWHILG, APGAPQTALLLPRGEDNHW PWNLGLCVLGSPRGLCPGH ACTCVLQVVSKYIESPVLEL DVGKVKFDIRYIVGGTSLAC PLKNPYVSAHLSSRAFAQG RGGPTWSGWLLARDPARVE LSSLVHSVERCPMQWALSE LHHLSTWSSGLCGLAG**VV LSSLVHSVERCPMQWALSE LHHLSTWSSGLCGLAG**VALCLLPGAADPLLRGC AS**LRJEPQAEIFRAFTELF AVMALCLLPGAADPLLRGC AS**LRJEPQAEIFRAFTELF KTGIQKYRREALLQHYLGP, AGRSKALTGTAERSCSSGS FTLCVNYNDSRAIMKTLAAI GGDCASKAEIQLVGUIGMPP FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIIS PPCHISVKSGAMLKASRLLLJ KEL 13907 44275 B 13987 1 20440		sequence		09/540,217		of peptide sequence	deletion, \-possible nucleotide insertion)
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PWNLGLCVLOSPRGLCPCH ACTCVLQVVSKYIESPVLFL							AGGPEGPPWWVLGWHHLGTQ
ACTCVLQVVSKYIESPVLFL	1		l		l	•	APGAPQTALLLPRGEDNHWICK
Dyckvkfdiryivgatslac Plknyyvsahlssrafaqgi RGGPTWSGWLLARDPARY* LSSLVHSVERCPMQWALSE LHHLSTWSGLGCLAG**VE HRTPTPP**QACSRPSGGL' A*WALCLIPGAADPLLRGC A*WALCLIPGAADPLLRGC A*WALCLIPGAADPLLRGC A*LRLPGPQAEIFRAFTELF CAS*LRLPGPQAEIFRAFTELF KTGIQKYRREALLQHYLGP, AGRSKALTGTAERSCSSGE FTLCVNYNDSRAIMKTLAA GFDCASKAEIQLVGILIGMPP FMYANSCKQMSQIKYTAN QMMIFGNEVKLMSVARTIS PCHISVKSGAMLKASRLLLI KEL 13907 44275 B 13987 1 2040 13908 44276 A 13988 205 438 LYFLETGVFCVQVPAQQC*C	1		l		1		PWNLGLCVLGSPRGLCPGHQR
PLKNPYVSAHLSSRAFAQGE RGGPTWSGWLLARDPARV* LSSLVHSVERCPMQWALSE LHHLSTWSSGLGGLAG**VF HRTPTPP**0,40ACSR*SGGL** HALSTWSSGLGGLAG**VF HRTPTPP**0,40ACSR*SGGL** A*WALCLLPGAADPLLRGC A*WALCLLPGAADPLLRGC AS**LR.PGPQAEIFRAFTELF GEGRQEAKKLKGSGQPLKT KTGQKYRREALLQHYLGP AGRSKALTGTRERCSSGSE FTLCVNYNDSRAIMKTLAAI GEPCASKAEIQLVOILGMP FMYANSCKQMSQIKYTAN QMMIFGNEVKLMSVARTIS PCHISVKSGAMLKASRLLJ KEL	1		1				ACTCVLQVVSKYIESPVLFLRE
RGGPTWSGWLLARDPARV-			1				DVGKVKFDIRYIVGGTSLAGTR
LSSLVHSVERCPMQWALSE LHHLSTWSSGLCGLAG**VF							PLKNPYVSAHLSSRAFAQGRRG
Lihilstwssglcqlag*-ve			1				RGGPTWSGWLLARDPARV*DL
HRTPTPP**QAACSRPSGGI- A*WALCLIPGAADPILGCC A5*LRLPGPQAEIFRAFTELF 13906			l				LSSLVHSVERCPMQWALSEWQ
A*WALCLLPGAADPLLRGC			l				LHHLSTWSSGLCGLAG**VFQM
A\$*LRJGPQAEJFRAFTELF 13906			ı				HRTPTPP**QAACSRPSGGLYR
13906 44274 A 13986 1 600 MEEQRVQFTLWLHFSPHGIC ESGRQEAKKLKGSGQPLKT KTGIQKYREALLOHYLOP, AGRSKALTGTAERSCSSGS FTLCVNYNDSRAIMSTLAA GFDCASKAELQVOILGMP FMYANSCKQMSQIKYTAN QMMIFGNEVKLMSVARTIIS P/CHISVKSGAMLKASRLLLJ KEL 13907 44275 B 13987 1 2040 13988 44276 A 13988 205 438 LYFLETGVFCVQVPAQQC*C			ı				A*WALCLLPGAADPLLRGCSEQ
ESGRQEAKKLKGSGQPLKT KTGIQKYRREALLQHYLGP, AGRSKALTGTAERSCSSGSI FTLCVNYNDSRAIMKTLAAI GFDCASKAEJQLVQILGMPP FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIIS PCCHISVKSGAMLKASRLLLJ KEL 13907 44275 B 13987 1 2040 13988 205 438 LYFLETGVFCVQVPAQQC*C	1		l				AS*LRLPGPQAEIFRAFTELFQV
KTGIQKYRREALLQHYLGP/ AGRSKALTGTAERSCSSGSL FTLCVNYNDSRAIMKTLAAI GFDCASKAEIQLVQILGMPP FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIB P/CHISVKSGAMLKASRLLLI KEL 13907 44275 B 13987 1 2040 13908 44276 A 13988 205 438 LYFLETGVFCVQVPAQQC*C	13906	44274	A	13986	1	600	MEEQRVQFTLWLHFSPHGICQF
AGRSKALTGTAERSCSSGSE FTLCVNYNDSRAIMSTLAA GFDCASKAELQLVQILGMP FFMYANSCKQMSQIKYTAN QMMIFGNEVKLMSVARTIIS PICHISVKSGAMLKASRLLLJ KEL							ESGRQEAKKLKGSGQPLKTGEI
FTLCVNYNDSRAIMKTLAAI GFDCASKAEIQLVQILGMPP FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIS PCHISVKSGAMLKASRLLLI KEL			l				KTGIQKYRREALLQHYLGPASL
GFDCASKAEIQLVQILGMPP FFMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIIS P/CHISVKSGAMLKASRLLLI KEL			l				AGRSKALTGTAERSCSSGSDSSI
FMYANSCKQMSQIKYTANN QMMIFGNEVKLMSVARTIIS PICHISVKSGAMLKASRLLLJ KEL			1				FTLCVNYNDSRAIMKTLAAIGT
QMMIFGNEVKLMSVARTHS PICHISVKSGAMLKASRLLLI KEL	1		l				GFDCASKAEIQLVQILGMPPER
P/CHISVKSGAMLKASRLLLI 13907			l				FMYANSCKQMSQIKYTANNGV
KEL 13907 44275 B 13987 1 2040 13908 44276 A 13988 205 438 LYFLETGVFCVQVPAQQC*C			1			1	QMMIFGNEVKLMSVARTHSKQ
13907 44275 B 13987 1 2040			1				P/CHISVKSGAMLKASRLLLE*A
13908 44276 A 13988 205 438 LYFLETGVFCVQVPAQQC*C							KEL
13300 11870 11 10300 100							
	13908	44276	Α	13988	205	438	LYFLETGVFCVQVPAQQC*CSV
			1				PGSCRRPLEEAGWRGLGLSGRL
	1						AGTGLPSGSRGNRTERGERGKL
VRGRRGVTRWLS							VRGRRGVTRWLS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino neid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13909	44277	A	13989	284	632	KTGVFCVPGPRPTVLM*CSFVG GKKNQYCLTGDAGCLQEKHSL GPAGV/HLEEAGWRGLGLSGRL AGTGLPSGSRGNRTERGERGKL VRGRRGVTRWLSHPQVQRQAD AGTCSPFEP
13910	44278	A	13990	594	1003	VRKGHHAVRGLCGRAQFRCIS GDPGKRPLG/LPGGNAPKSGSL ARRPQPVRIPGSRPQ/RGSVSHL GPAGVH*R/RAGWRGLGLSGRL AGTGLPSGSRGNRTERGERGKL VRGRRGVTRWLSHPQVQRQAD AGTCSPFEP
13911	44279	A	13991		1500	VRVAHIPIAAAEIAAIPDRGGHL SVGKRKLTTWFFYGVLPFRDV AIEFSPEEWECLDSAQOHLHRD AMLENYGNLVSLAAGDQSLPG DAAASAVCRQLSPSAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGALIG PPVNGVJSPAGGAGLARVTVSQ KCDPRHGGGCALALQGWTLTVP SESPSRESKE LGGL WSFRADLD SRRQGFEEQWYPRPLRELRGRG GRRPEARRWEGGSPGCKLLPI SRPELSPKRLVNGLRGAGL GDGETGGTGGGGGRRKKRC TLSRDEGFNDICQDWLRCQF VGWVLYEQEVTLPEQWTQHLR TRVVLRIASAHSYATVVSAARS QVFEPHAAGYVRECLIAGPLLM PYTTCCPLLISPVSAVGEWGT RA*GADSSSLVQNTDFDFFRNYA GLQRSVLLYTTLTTYDDITITT GLYPHDSABIPFPFYLYNTGN
13912	44280	Α	13992	2	307	
13913	44281	A	13993	220	427	MSTLPFPSLQMFRRPPYGSPSTC RSQDNRSYSRVTEELQERRRW NSGDAN*DQSPEARRQRPRHIE EAA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13914	44282	Α	13994	1	2832	MGKYYTQGDKVLMPLAIQVH
		l				HAVCDGFHVGRMLNELQQYC
	1	l				DEWQGGAYPKGYFVQNTDFDF
		l		1	l	FNYAGLQRSVLLYTTPTTYIDDI
		l		1		TITTGVEHDSAAAPMVDSLIAR
	l	l		1		VGVMARGNAITLPVCGRDVKF
		l				TLEVLRGDSVEKTSRVWSGNE
		l				RDQELLTEDALDDLIPSFLLTGQ
		l				QTPAFGRRVSGVIEIADGSRRR
			ļ			KAAALTESDYRVLVGELDDEQ
		1				MAALSRLGNDYRPTSAYERGQ
	1	1				RYASRLQNEFAGNISALADA
13915	44283	Α	13995	1	2934	MITFDFMSHIQVTLMQEVSFHG
						LGQLRPCGFAGYSPPPGCFHRL
						ASSVCSFSRCTVQAAVDLPFWG
						LEGTSPLLAASLGSAPVGTLCG
				l		GSDPTFPSWTALADVLHEEGHT
		1				PASNFCLRNLAGEASQSWRKE
		1				GLGTPRPTASSTEELQTLNEDSR
	İ	l		l .		LMTSPHNLEETRDCDTELPKKS
	1	ŀ		l		LGWKEIRVYGQNKTLGFWLQA
						QLLICDGGNLRMFRLLPYGSPS
				ŀ		TCRSQDNRGYGGVTEELMERK
						RWNSTDGFHIQGVLE
13916	44284	Α	13996	68	425	SHILPGAPGAPAWWTRWPSTLP
				1		EPFPRGRGSPAGTSPISRPGLVQ
l						SS*ASRGSDSRLPV/GPASCQAS
Į.						GPGPDSRRPPPCTPA\GPHHGSL
				ľ		PSAGRVGASAAAAGPPSPAVPL
						PPAERPAP
13917	44285	Α	13997	324	723	TAFSLNFQGGVEGEVWAGTRA
						ARGACAGERGLGGSRTRSGW/P
						GTAGPGQAGHGTCSLPCPSLPP
1						AMGSCVSEPPRQAPPPAPWRLV
l		1		1		PSTAQGLRSVGTQCGTGGSSAC
1		l	l	1		GPC\GDPGF*THQSAPCQNGPIS
				·		SL
13918	44286	Α	14000	39	383	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	i .	1		sequence		·
		<u></u>				
13919	44287	Α	14001	181	1446	EDLLQEAQALAEVTAQGKGLG
		1		ł		QVRSLDLSLEACAGSPPLQVLD
						LQDREPRSLTSLPSSAANRGKL
l		1				EKAQSNPCGHRRHQAAGLKLP
						LFSAQAPQGQPTPAPRFRAATR
		1				KPGTAQSHAPLWSPAMSFFPEL
					l	YFNVDNGYLEGLVRGLK/ALGE
	İ	1				LSQADYLNLVQCETLEDLKLHL
						QSTDYGNFLANEASPLTVSV\ID
		1		l		DRLKE\KMV\VEFRPHEGTMPY
						EPTSPSFP*TSITLQFT*FDNVI/L
1	1	1			i e	WLIHRARLHORSIAELVPOVAH
						PLG\SFEQMEAVNIAQTPAELY
	ŀ					NAILVDTPLAAFFQDCISEQDLD
					i	EMNIEHRNTLYKAYLESFYKFC
1						TLLGGTTADAMCPI\LEFEA\NR
l	İ			l		RAFINT\INFFG\TKTVPKKDRCP
						AFFPH/CGAGFFPEGPAAMGFG
					i	LNNLLTGAKKRGPYNFPEVOL
ĺ		l				AFLRGDG
13920	44288	A	14002	1	1182	
13921	44289	A	14003	26	342	ARFLAGP*LPSRRTGLRDLOPA
						MPEPP\PTPWAPVRPEPPRRAPP
		l			1	PAPRRPVPSTGKEQTTQGLRSA
1	1	ı		i		GAGHRDWQAAPPAAPVRDPLG
	1					EASWAPESGGDVESLYV
13922	44290	Α	14004	66	463	
13923	44291	A	14005	3	1446	
13924	44292	A	14006	221	852	NSFLMVGFFPFFSL/CAAFDTKT
						GLRVAVKKLSRPFLRPDGYYTE
				l		FNIYLLFNFYILTSSAV*VVLSH
						SVVIECLLSIRHCSRYFRKSSED
				l		KNFLRGSRLLCNFSLISRYLVTH
						LMGADLNNIV*RYR*YK**FFK
	l			l	1	MNSPFLPLPVGGELIMHHKVDP
	ŀ					VCFVFFSLPVSKS*QFVCSSSAL
						*ILDFGLARHTDDEMTAKGOPE
				l		CFVRWACAS
13925	44293	A	14007	55	784	RHIQDPASQRLTWNKSPKSVLV
	1	1		1		IKKMRDASLLOPFKELCTHLME
	l			1		ARGAGGGHDARSLTTAPVSRE
	1	1				PPCRRVGSAGGMSRLALCFQEN
	1	1				MIVYVEKKVLEDPAIASDESFG
	ľ	1				AVKKKFCTFREDYDDISNQIDFI
	1					ICLGGDGTLLYASSLF/QGNAA
	ł			l		VVLRSRLKVRVVKELRGKKTA
	[VHNGLGEKGSQAAGLDMDVG
	1			I		KQAMQYQVLNEVVIDRGPSSY
		1		I		LSNVDVYLDGHLITTVQGDGVI
				I		VSTPTG
		Ь_				101110

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion)
13926	44294	A	14008	155	1592	PESRIQCFSDRSANFI.TMEMEQ EKMTMKELSPDAAAYCCSAC EKMTMKELSPDAAAYCCSAC HOGDETWSYNPIPRGRAKSRSL SASPALGSTKEIRKTRSLHGOPC YTTFOFRACVLQNPQTIMHIOD PASQRIHVEQRPPRTVLVIKKM RDASILLOPFKELCTHLMEENMI VYYEKKVLEDPAHASDESSGA VKKKFCTFREDYDDISNQIDFII CLGVDGTILLYASSLFPGQPSP VMAFHLGSLGFLYTBISFENFOS QVTQYIEGNASCCSSGSRLKGQ GW*RSLRGKKTAIVHNGLGEK GISQACRPGTMDVGKA/QAMQ VPKS*NFEVWIDRGPSSYLSNV VYLDGHLIITVQGDGKARSTV
13927	44295	A	14009	327	466	LGP*ASLGRE*RLRLSLSGVIVS TPTGSTAYAAAAGASMIHPNVP AIMITPISRSVLPTAGGPHLSLL DPQIMLSPEARNAGGGPFWKP WLRLLLSRPSISITTSCYPLPSIC VRDPVSDWFESLAQCLHWN
13927	44295	A	14010	2	196	
13929	44297	Α	14011	923	1776	WRKCGCETAGSASATRAAATE KMEAPRNKMCRDQENSAWRN AVPGPE*ROPEEPPRTTSILAEP TVLSRRKCRPSEKRGSATEKFG ATSAVTENPPLGAAGERANKTL GAATVPPTLÖPRPMEKGGTR CPSTL*PSVQERQPESPSGNIPJE* TOKIEFAGRSVTFQHRWATTLR TRISGTGASPRORSAFFGFPWC G/AAQPSDEESTPATQEEAQAQ G/AAQASSEEPGHCAPRPPHA PRSALFEDWSREPERSDDGDLS ECRRPATEPWRPPGSAWRLPCA A
13930	44298	Α	14012	568	1223	SARGILPGSOPLSHAGPOLOVE WPGCLPPPDSTILELVVCGTVAI SRVRMSAVCVLRTWS\RNAGQ LICETVFQPCSNVIHVLKGPNY VCFGAYPSRVSHPHNFLKTI NAVCGQLVQFRPPDTEEGIRK VQSDKASVTNASPYDGV\RKL* HSLDDIAYYGKPLVNIETEALK GTVNLFYQIDYCPPFVIGSQLK

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		1		sequence		
						<u> </u>
13931	44299	A	14013	5	296	LFSRVPSKGPPRYHHTSSRG*TR
					1	AD*LISS\GNS*AHCIPSGLQGM
		ı				LNLLPTPRAPVRPEPPRRAPPPA
						PRRPVPFDHPRACGTGRQLHLQ
						PRCGIH
13932	44300	Α	14014	1	376	MQAVLLTVLQKHNQSVIYWY
						MKYIYYEGLVIMDDYDYVIMD
l l						YMTMETEKSHDLQAGGPGKPV
						VQFQFKPKILQASRGNGVKSWS
l		1				KSKGLRTRCANVQGQENREGP
l		1		l		AKHSPFLCLFVLFGSSTGRMIPN
						HTGDTSKVIAGTEERGVPKKVT
		1				LKFDACAAIDRPWAHQLDCTY
1						HFKGSNDEHDSRLHTNAG*L*L
		1				C/RLWIT*LWRLRNPMICRLEA
1		1				QESOWYSFSSSPKSCKPVEAMA
						LSPGPSPKA*EPGVLMSKGRKI
		1				GRAQLSIHLSSAFLFFSAPOOVE
13933	44301	A	14015	1638	2289	VEKKKPMSAPALRLPDLTKLF
13933	44301	n	14013	1030	2209	TLYVSETKKMAVRVLTQTVGP
		1			i	WPRPVAL\LSKOLDEVSKRWPP
					i	CPKSLVAIALLAOEADKLTLRO
Į.					i	NLNIKSPCAVVILINTKGHH*LM
		1				
1						NARLARYQSLLCEHPRITLGSL
		l				QHLNPATFLPVSESPVKHNCVE
						VLDSVYSYVGPNHRDHPLNISR
		1				LGAVTWMGSSFHQPLQSDTLK
		_				KTTSPGSSHTRKLTGPRTAEA
13934	44302	A	14016	1	675	MAPPLRPLARLRPPGMLLRALL
	1					LLLLLSPLPGLREGIGELITPIGT
1	1	1		1	İ	SLPDLDPARRRWEGGIGRVGSE
1						VADLCPGKEGGKVPEAEKEGV
l			1		İ	WCFSELSFVKEPQDVTVTRKDP
ŀ		1	İ	l		VVLDCQAHGEVPIKVTWLKNG
ŀ					1	AKMSENKRIEVLSNGSLYISEV
1						EGRRGEQSDEGFYQCLAMNK\F
1	ł					*AILNQKAHLALSRIGST*RRRP
						DRP*EDEAFVMTTHCFQDLLTS
13935	44303	C	14017	18	329	
13936	44304	Α	14018	391	895	WASSDTPTGFTQRWRPDWAGG
		1				GGGEERGDGTGVVWEGEKVY
1	1	1	1			RFRRVTGGSSFPSPAVLVRAPN
					İ	TGMPKLTYIQT*GPLGERGKEP
1	1	1	1	1	1	VPPN\PTLPQTFGGLLNASVVRK
	1	1		l		TEIPSPSPPPVCYLRASAERO/PA
1	1	1	1		1	GGSPOAKWRRAASSPPPLPPVH
1		1			1	LPPPPPPPRPPPPPPP
13937	44305	В	14019	1	1110	
13731	177505	10	1.7017	ı·	1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13938	44306	Α	14020	282	710	YPEPQGWRVAHTVQYPIHIQPL
		ı				KPGPRGR*SIMDIQVPPKEGGV
		l				LILKN\SSRLLNSWTQGCLQISA
		l				FSPTSSTLPGTPWSNVGPYKIAP
		l		ł		ITSFGE\PFIRVIWKQLPSWISIIQ
		l				V/SHKQ*QIHRSVDIGWSPGQEC
	i	ı				GPKGCFV
13939	44307	A	14021	1	1416	MIILIDAEKAFDKIQQPFMLKTL
		l				SKLGTDGTYLKIIRAIYDKPTAN
		l				IILNGQKLEAFPLKTGTRQGCPL
	İ	l				SPLLFNIGLEDLARAIRQEKEIK
		l		l		GIQLGKEEVKLSLFADDMIVYL
		ı				ENPIVSAQNLLKLISNFSKVSGY
		l		ŀ		KINVQKSQAFLYTNNRQTESQI
		l				MSELPFTIASKRIKYLGIQLTRD
		l				VKDLFKENYKPLLKEIKEDTNK
		ı				WKNIPCSWVGRINIVKMAILP\R
		l		ł		DIDQW/NRTEPSEIMPHTYNYLI
		l				FDKPEKNKQWGKDSLFHKWC
		ı				WENWLAVCRKLKLDPFLTPYT
		l		İ		KINSRWIKDLNIRPKTIKTLEEN
		l				LGITIQDIGVGKDFMSKAPKAM
		l		ŀ		ATKAKIDKWDLIKLKSFCTAKE
		l				TTIRVNRQPTTWEKIFATYSSD
		l				KGLISGIYNELKQIYKKKTNNPI
		l				KKWAKDMNRHFSKEDIHAAK
		l		l		KHMKKCSSSLAIREMDIKTTMR
		I				YHLTPVRMAIIKKSGNNRCWR
		L				GCGEIGTL
13940	44308	Α	14022	1	3450	
13941	44309	В	14023	1	3171	

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13942	44310	А	14024	1	2433	MRTKTQHTRISGTHSKPVCRGK FIALNAHKRKQEKSKIDTLTSQ LKELEKQEQTHSKASRROEITKI
		1		ł		RAELKEIDTQKTLQKINESRSW FFERINKIDRPLARLIKKKREKN
	ŀ	1				OTDTIKNCKGDITTDPTEIOTTI
						REYYKHLYANKLENLEEMDKF
						LNTYTLPRLNQEEVESLNRPITG
	1					AEIVAIISSLPTKESRTGWIHSRI
						LPEVOGGTEKEGILPNSFYEASI
	1					LIPKPGRDATKKENFRPISLMNI
						DAKILNKILAKRIQQHIKKLIHH
	1					DQVGFIPGMQGWFNIRKSINVI
						OHINRTKOKNHMIISIDAEKAF
	İ					DKIOORFLLKTLNKLGIDGTYF
		l				KIIRAIYDKPTANIILNGQKLEAF
		l				PLKTGTROGCPLSPLLFNIVLEV
		l		1		LARAIRQEKEIKGIQLGKEEVKL
		1				SLFADDMIVYLENPIVSAQNLL
		1				KLISNFSKVSGYKINVQKSQAF
		l				LYTNNTQTESQIMSELPFTIASK
		l				RIKYLGIQLTRDVKDLFKENYK
		l				PLLKEIKDDTNKWKNIPCSWVG
		l				RINIVKMAILPKNWKKTTLKFI
	ŀ	l	i	l		WNQKRAHITKSILSQKNKAGGI
		l		l		TLPDFKLYYKATVTKTAWYW
		1				YQNRDIDQWNRTEPSEIMLHTY
		1		i		NYLIFDKPEKNKQWGKDSLFN
		1				KWCWENWLAICRKLKLDPFLT
						PYTKINSKWIKDLNVRPKTIKTL
		1		l		EENLGITIQDIGMGKDFMSKTP
						KAMATKAKIDKWDLIKLKSFC
13943	44311	Α	14025	1	1023	MGAIYDKPTANIILNGQKLEAF
		1				PLKTGIRQGCPLSLLLFNIVLEV
		l				LARAIRQEKEIKVIQVGKEEVK
						LSLFADDMIVYLEDPIISAPNLL
						KLISNFSKVSGYKINVQKSQAF
		1		l		LYTNNRQTESQIMSELPFTIASK
						RIKYLGIQLTRDVKDLFKENYK
		1		l		PLLKE\KEDTNKWKNIPCSWVG
		1				RINIVKMAILP/KAICRKLKLDPF
			1	l		LTPYTKINSRWIKDLNVRPKTIK
						TLEENLGNTIQDIGMDKDFTSK
				l		TPKAMATKAKIDKWDIIKLKSF
						CTAKETTIRVNRQPTEWEKIFAI
	1			I		YSSDKGLISRIYNELKQIYKKKS
			l	1		NNPIKKWAKDM/NRHFSKEDIY
		<u> </u>				AAKRHM
13944	44312	В	14026	1	2206	
13945	44313	Α	14027	1	1689	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13946	44314	A	14028	11	2523	MKQFLLYLDESNALGKKFIIQD
13340	44314	ı^	14020	ľ	2323	DDTHVFVIAELVNVLQERCHTR
1		l				LGYTEFLVAWRVTFGLCVEAV
1		l				TLHLKYOILIRGLLEMMSFSDA
1						DILKQLPVTVPGLFPASLSPSSL
1						LGNSPPSWLRHNSESKVSAVSS
						PSATKTLSTGIGKLDPGHKEMA
		i i				EESELLKNKMQAPPLSRCPESQ
}		i i		l		KCQHQLRLHHWKPSVRHQVKR
		1				RSPAVLRSAMPPADCPAVLEAT
		l				TATHPEKGTALSKHLPSSDSMS
		1				LKVDVEALENSPGATYIWKGG
						KVTRDSQPKEQGKGDLKKKKK
						GKLPKNYDPKLTPDPERWLPM
1						QECSFYQGRKKGKKKDQMGK
1		1			i	GTQGATAGASSELDARKTVSSP
		1				PTSPRPGSAATLSASTSNIIPPRH
1						QRPAGAPATKKKQQQKKKKG
						GKGFPVLREITVVKVDTLVVFQ
1	İ	l				ILEERLSVFHIQYDTSYPFSTVDI
Ì		1				EDHECAVWLLLRKSKSDDKTT
		l				RLEAVREMSETHHWHDAEKAF
		l				DKIQQPFMLKTLNKFGVDGTY
		1				LKIIRAIYDKPTANIILNGQKLE
		1				AFPLKTGTRQGCPLSPLLFNTV
		l				LEVLARAIRQEKEIKGIQLGKEE
		1				VKLSLFAGDIIVYIENSIVSAPKL
		l				LKLISNFSKVSEYKINVOKSOAF
		ı				LYTNNRHTESQIMSKLPFTIATK
		i				RIKYLGIQLTRDVKDLFKENYK
		l			l I	PLLNEIKEDTNEWKNIPCSWVG
		1				RINIMKMAILPKVIYRFNAISIKL
13947	44315	A	14029	I	2868	
13948	44316	A	14030	I	3099	MGELITPLSTLDRSTROKVNKD
		ŀ				TOELNSALHOGDLIDIYRTLHP
	1					KSTEYTFFSAPHHTYSKIDHILG
						SKALLSKCKRTEIITNYLSDHSA
1						IKLELRIKNLTONRSTTWKLNN
						LLLNDYWIHNEMKAEIKMFFET
ł			i			NENKOTTYONLWDAFKAVCR
1						GKFIALNAHKRKQERSKIDTLT
				1		
1	1	1	l	1		SQLKELEKQEQTHSKASRRQEI
1		1	l	1		TKIRAELKEIETQKTLQKINESR
1		1	l	l		SWFFERINKIDRPLARLIKKKRE
-		L.				KNQIDTIKNDK
13949	44317	A_	14031	2	3419	

SEQ ID SI	EQ ID NO: [Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO: of	f peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon. /=possible nucleotide
se	equence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13950 4	4318	A	14032	I	2685	MGDFNTPLSTLDRSTROKVNK
13930 4	4318	^	14032	1	2003	DTQELNSAPHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKTDHIL
						GSKALLSECKRTEITNYLSDDS
	1					AIKLELRIKNLTQNRSTTWKLN
						NLLLDDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFVALNAHKRKQGRSKIDT
						LTSQLKELEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTVQKINES
	- 1					RSWFFERINKIDRQLARLIKKKR
						EKNLIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
				1		DKFLDTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLTTKKSPGPDG
						FTAEFYQRAIRQEKEIKGIQLGK
						EEVKLSLFADDMIVYLENPIVS
	1					AQKLISNFSKVSGYKINVQKSQ
						AFLYTNNRQTESQIMSELPFTIA
						SKRIKYLGIOLTRDVKDLFKEN
				ŀ		YKPLLKEIKEDTNKWKNIPCSW
				ŀ		VGRINIVKMAILPKVIYRFNAIPI
						KLPMTFFTELKKTTLNFIWNOK
				1		\RAHIAKS/VLSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYON
1 1						RDTDQWNRTEPSEIMPRIYNYL
						IFDKPEKNKQWGKDSLFNKWC
	- 1					WKNWLAICRKLKLDPFLTPYT
						KINSRWIKDLNIRPKTIKTLEEN
	- 1					LGITIQDIGMGKDFMSKTPKAM
	1					ATKAKIDKWDLIKLKSFCTAKE
						TTNRVNRQPTKWEKIFATYSSD
13951 4	4319	A	14033	845	1616	ARAEVKLSLFADDMIVYLENPII
13931 4	4319	^	14033	043	1010	*ARAEVKLSLFADDMIVYLENP
						IISAQNLLKLISKFSKVSRYKINV
						OKSOAFLYTNNROTESOIMSEL
						PFTIATKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWIGRINIVKMAILPKVIYRF
						SAIPIKLPMTFFTELEKKNWLAI
						CRKLKLDFFFIPYTKINSRWIKD
						LNVRPKTMKTLEESLGNTIQDI
						GIGKDFMTKTPKAMATKA/KKS
						FCTAKETTIRVNRQPTEWEKIF
						AIYPSDKGLIS

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon,/=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ĺ	sequence		
13952	44320	Α	14034	707	2878	TLMQKSSIKYWQNESSSTSKSL
13752	77520	^	14054	1,0,	10,0	STMIKWASSLGCKAWFNIRKSI
1				1		KVIQHINRAKDKNHMIISIDAEK
1			l			AFDKIQQPFMLKTLNKLGIDGT
l						YFKIIRAIYDKPTANIILNGQKLE
i						AFPLKTGTROGCPLSPLLFNIVL
						EVLARAIRQEKEIKGIQLGKEEN
						KLSLFADDMIVYLENPIVSAON
1						,
						LLKLIRNFSKVSGYKINVQES\Q
						AFLYTINRQTESQIMSALPLTIA
						SKRIKYLGIQLTRDVKDLFKEN
						YKP/LLKEIKEDTNK\WKNIPCS
		1				G\EGRINIVKMAILPKNWKKTTI
		1	i			NRFIWNHKRACIAKTILSQKNKA
					ŀ	GGITLPDFKLYYKATVTKTAW
						YWYQNIDIDQWNRTEPSKIIPRI
						YNNLIFDKPDKNKKWGKYSLF
						NKWCLENWLAICRKLKLDPFL
						TSYTKINSRWIKDLNVRPKTIKT
					-	LEGNLGNTIQDIGMGKDFMSKT
	İ					PKAMATKAKIDKWDLNNLKSF
						CTAKETTIRVNMQPTEWEKIFA
						IYPSDKGLISRIYNELKQIYKKK
			l			TT\NPIKKWAKDMNRHFSKEDI
						YAAKKHMKKCSSSLAIREMQI
				ł		KTTMRYHLTPVRMAIIKKSGNN
						RCWRGCGEIGTLLHCWLDCKL
					1	VQPLWKSVW*FLRNLELEIPFD
				1		PAIPLLGIYPNDYKSCCYKDTCT
		Ì		1		RMFIAALFTIAKTWNQPKCPTII
		l				DWIKKMWHIYTMEYYAAIKNE
1	1	1				EFVSFVGTWMKLEIIILSKLSQE
13953	44321	A	14035	869	3869	RHKKPFKKLTNPGAEIOTTIREY
		1				YKHLYANKLENLEEMDKFLNT
1						YTLPRLNQEEVESLNRTITGSEI
	ł					EARINSLPTKKSPGPDGFTAEFY
						QRYKEEMVPFLLKLFQSIEKEGI
1						LPNSFYEASIILIPKPGRDTTKKE
	ŀ					NFRPISLMNIDAKILNKILANOI
						OOHIKNLIHHDOVGFIPGMOG
			l	I		WFNICKSVNVIQHINRTKDKNH
				l		MIFSIDAEKAFDKFQQLFMLKT
		1	1	I		LNKLGIDGMYLKIIRAIYDKLT
			1	l		ANIILNGQ
12054	44322		14036	2	99	MILINUQ
13954	44322	Α	14036	<u> -</u>	77	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	endon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
13955	44323	Α	14037	1	914	TVTVIHTVARVGNLDIQIQRKS
13933	44323	^	14037	ľ		ELFLSLVVCEAIKNEMNVMLKS
	i					SPVDLVTATDOKVEKMLISSIK
1						EKYPSHSFIGEESVAAGEKSILT
		ŀ	!			
		ŀ	1			DNPTWIIDPIDGTTNFVHRFPFV
1		ŀ				AVSIGFAVNKKKKVDETYLGL
		l				VKETCFPLQIEFGVVYSCVEGK
						MYTARKGKGAFCNGQKLQVS
1						QQEDITKSLLVTELGSSRTPETV
1	l			i	1	RMVLSNMEKLFCIPVHG\IRSVG
	l					TAAVNMCLVATG\GA\DAYYE
1		ľ	ŀ			MGIHCW\DVAGAGIIVTEAGGV
i	ŀ	ļ				LMDVTGGPFDLMSRRVIAANN
						RILAERIAKEIQVIPLQRDDED
13956	44324	В	14038	91	1959	
13957	44325	Α	14039	28	152	LPSPSSKDTDS/AASCCAPRGRA
		1				KGSVSNSRLSFPPSSVPLP
13958	44326	A	14040	35	879	GSQCSSPPPASRCGSCSSAPSRH
						CPCRDFEPAVAGLARTPVLTTK
	i					TNPSPPARTGLNAPSMGTSLI/P
						ALCCFPLYQGSTELQCKAP**LP
			Ì			SPSSKDTDS/AASCCAPYFLVLL
						KVLSSVDNSSIWCSAGPPAQS/S
					i	SLYTIPMSSMSAKNCSVLRSGT
		l				WASFTCS*GSQRVSLGTPEALG
						TEDHTSRSLGHSHVCHVDLSPA
						TASCSFQTASSEVSPPATASCSF
						OTASSEVSPPATASCSFOTASSE
						VSPAATASCSFTFSFFFVCALWF
						GICPSCSAAVLST
13959	44327	A	14041	3	159	EHQCENPQ*NTGTPNPAAHQK
1.5555	1.1227	ľ.				AYPP*SSGLHPWDNQRQKPHD
						YLNRCRKGL
13960	44328	A	14042	54	245	GGGGGEAEDRLREPETEKALSS
1.5500	1	ļ.,	1	ľ.	I	SL/RRAPDQC/RALIMQLFQAHC
		l		1		FFLST/QATAALQAHYAHIFPSK
13961	44329	A	14043	903	1205	WDVSONNKSYL*OTHSOYHTE
13901	77329	ľ	17073	"	1203	WAKTGSIPFENWHKTGMPSLT
		l		1		TPIQHSVGSSGQGNQTNREPNH
		İ				E*TPIHNCFKENKIPRNPTYKGC
1		l	l	1		
100 (0	11000	_	14044		220	EGPLQGELQTTAQ
13962	44330	В	14044	I	339	
13963	44331	В	14045	1	1547	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13964	44332	A	14046	807	1440	NTGKPNPAAHRKAYPPRSSRLH PPDLQQIPTDLQLDITGRKT NKQKGIASPSTKRTSTFKPHLQ ANAQRICHHQACVTRVPKGST KHGKEQL VPATAKTCQIVKNIN AMKKLHQLTEIQITIREYHKHL YANKLKNLEEMDKFLDTYTLP RLKQEEVESINPIMGSEIEAIN SLPTKKSPGPDGFTAKFYERYK EEL VSFLLKLFQSIEKEGILPNSF YEASIILIPKPGRDTTKKENFRPI SLKNIDAKILNKILANRIQPIEK LIHHDQVGFPEMKGWFKKCKS INVIHHINQTNDKNHMIISIDAE KAFDKTQQFPMLKTLNKLGTIR GKERIKGIQGKEEVKLSLFAD DMIVYLENSIVSAQNLLKLISNF SKYSRYKINLQKSQAFLYTNNR HMESQITSELPFTIDTKRIKYLGI QFTRDVKDLFKENYKLLLNEIK
13965	44333	A	14047	603	764	LPLFLIEYPLFLSLA*LPWPKLPT LC*IGVVREGIPVLCCFSKGMLP VFAHSV
13966	44334	A	14048	1566	1865	MOSIRKTIGIHAKYDKACRNGE DICLRETNSAADTQAKRV*SGP PANSNRPAAEGPDC*KEN*QTE RTSTPKPHLYVTIIKDQRGTQLL TNNGTKLDRE
13967	44335	В	14049	1	1932	
13968	44336	A	14050	711	1575	TRKFRNG*IP*HIHPKTKSGR SRIFNRPITGSEIEAINSLPTKKS POPDGFTAEFYORYKEELTNIL DEHRCKILNKILANGIOHDIKKI INDOVGFIEPMGGWENTIKSI NVIQHYNRTROKNIHMISIDAE KAFDKIQOFFMLKTINKLGIDG MYLKIIRAMYDKPTANILINGQ RLEAFPLKTGTRQGCPLSPLLFN VILEVLARAIGQEKEIKGIOLGK EEVKLSLSADDMIVYLEKPIISA QNLFKLISNFSKVSDTKSMYKN HKHSYTPITDKQRAKS

1727

SEQ ID NO:			SEQ ID NO: in USSN	Nuclcotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	nou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence			sequence	,-,	
13969	44337	A	14051	i	1579	L MQGWFNVRKSLNIIHHINRNND
		1				KNHIIISIDAEKAFDKIQQPFML
						KTLNELGIDGTYLNIIRTIYDRP
				:		AANIILNVQKLEAFPLKTGTRQ
						GCPLLPLLFNIMLEVLARAIRQE
						KEMKGIQLGKEEIKFSLFADDII
1						VYLENPIFSAPNFLKLISNLSKFS
1						GYKINVQKSQAFLYIINRQTESQ
						IMSELRFTIATKRIKYLGIHLTR
						DVKDLFNENYKPLLNEIKEDTN
ł						KWKNIPCSWTGSINIVKTAILPK
				1		VIYRFNAIPIKLQLTFFTELEKTA
			İ	1		LKFMWIQGYSNQNSMVLTLLV
						IPRKMRSAVELQQTSTDLQLRD
			l	i		LTVRKKTNKRKATASTSQKGH
						LHQNPICRSPTSKTKEVGFRRSV
			l			ITNFSELKEDVRTYHKEAKNLE
						NRLDEWLTRINSVEKTLNDLKE
			Ì			LKTMARDLHDACTSFNRRFDQ
						VEERVTVTEDQINEINDGENGT
	i					KLENTPQDIIQENFPNLA\SRPTF
						KFRKYREHYKDTPREEQPQDT*
			i			LSDSPRLK*RKKC*RQPERKVE
						LPTKGSPSD
13970	44338	A	14052	50	1105	TRKSRRNG*IPGHIYPPKTK\QE
			l			EGESLNRPITGSEIEAIINNLPTK
			l			KSPGPDGFTAKFYQSTNNKNH
						MIISIDAEKAFHKIQQPFMLKTL
						NKLGIDGTYLKIIRAIYDKPTAN
1						IILNGQKLEAFPLKTGTRQGCPL
						SPLPFNIVLEVLARAIRQEKEIN
l						CIQLHNEEVKLSPFADDMIAYL
1			İ			ENPIVSAQNLLKLISNFSKVSGY KINVQKSQAFLYTINRQTESQI
ł						MSELPFTIATKRIKYLGIQLTRD
1		l	i			VKYLFKENYKPLLNEIKEDTNK
			ŀ			WKYIPCSWIGRINIVKMAILPKV
						IYRFNAIPIKLPMTLSSQNWKKT
						TLKFIWNOKRARIAKSILSKKN
						KAGSIMLPDFKL
13971	44339	A	14053	l .	764	MQGWFNVRKSLNIIHHINRNND
133771	111337	ľ.	11000	[/°•	KNHIIISIDAEKAFDKIQQPFML
			1	Ì		KTLNELGIDGTYLNIIRTIYDRP
			l			AANIILNVQKLEAFPLKTGTRQ
			1			GCPLLPLLFNIMLEVLARAIRQE
1			1			KEMKGIOLGKEEIKFSLFADDII
						VYLENPIFSAPNFLKLISNLSKFS
1			1			GYKI*FPLFLIEYPSFLSLA*LPW
						PALPTIC
L				L	L	1 15

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12072	14240	+-	14064	li i	344	MCDIODDIVA PAISVAIOS A CODEV
13972	44340	A	14054	l'	344	MGINQRRKAENSKNQSASSPTK
						DCSSSPAKEQSWTENDFEELTE
1		1			i	VGFRRQANAERFCHHQVCPKR
1						APEESTKHGKKQ/RSISHCKNM
					ļ.	PNCKDHRCYEETASTDSLAETQ
		<u> </u>				QQKREF
13973	44341	A	14055	301	370	DDGVF*IDNHVICKQRQFDFLFS
13974	44342	A	14056	1	1418	MVREGFPEEVTFELKNELEISK
		1				GQQMGKNIPDTGASSAKALGE
					ŀ	RELGTLDKMKNSPSDGSKKDIR
		1			1	THRKEAKNLEKRLDEWLTRINS
				i .		VEKTLNDLMQLKTMARELHDA
l				1		CTSFNSQFDQVEERINKIDVRLA
					1	RLIKKKREKNQIDAIKNDKGDIS
1	ł	1		i		TDPTEIQTTIREYYKHLYANKL
]		1	1	ł	ENLEEMDKFLDTYTLPRLNQEE
				1		FESLNRPIIGSEIEAVSNSLPTKK
			İ			SPGPDGFTAEFYQRYKEELVPF
	İ			l		LLKLFQSIEKEGILPNAFYEASII
				i		LIPKPGRDTTKKENFRPVSLMNI
						DAKILNKILANRIQQHIKKFIHH
	Ì					DOVSFIPRMOGWFNICKSINIIH
						HINRTNDKNHMIISIDAEKAFDR
				l		IQQPFMLNTLNKLGIDGMYLKII
				1		RTIYDKPTANIILNGQKLEAFPL
						KTGTRQGCPLS/ATPVQHSVGS
			ĺ			SDQGSQARERNKGYSIRK*GSQ
						IVSVCR*HDCIFRKPHRLSPKSP
13975	44343	Α	14057	1240	1596	CYSFLFVRCPSDSQVPQLQVC/C
						EFAGGPLQTVFAWVSPAEVVE
						MFLQG*RKPFYMSY/TSFEMAL
					İ	MKVWTMVRTSASPVGDVFGR
						GWSPFPTSTVGALTVFGVSPGS
				l		CRSSPLPSERLWV
13976	44344	A	14058	507	971	PANQKKSRTRRIHSQILPEAQR
						GAGTIPSETIPINGKRGNPPKLIL
						*SQHHPDTKAWQRHNRKREFY
					1	TDIPDDH*CKNPQ*NTGKPNPA
	1	1		I		AHQKAYPPRS\$WLHLWDARLV
1	1	1	I	I		QRTQISKRNSSYKQNQRQKPHD
						YLSRCRKGLQQNSAALHAKNS
13977	44345	la	14059	561	758	
13978	44346	A	14060	2364	2663	DGQLTLVSILNHYSLAQSCEY*
1.57,0	. ,540	Ι΄.		[· · ·		DGECFCLHCSASPFL*EGAOST
1		1				ACTAPPECFASDLGAV*LPQHS
1		1	l	I		WCSTLLLEGAPWPGTLGTAPOF
1		l		l		QHRKLGTNLAG
	L	L	L		L	Kukkrollarvo

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hed	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>				Language Communication in the communication in the
13979	44347	Α	14061]1	64	MPIGKVGEYYIKGTLRATKESE
		l			ľ	QQSSALELSSDRVYLNEKEPGD
		l				QPWNAAFVFRGCGVALALRPP
		l				HLGWGEPRWAGFRGAFGFPFS
		l				TLNPSNWSLTLIIQGCKQEGFYS
		l				PSMRPRVVPKRGSAPKVRAEEN
	1					AALPSRCPPGPLPVAQP/TGPAD
1 1		l				GGSCEQCPSCGCGCCQLPPPLP
		l				WFPR*TPCDKRV
13980	44348	A	14062	1	774	MFVWSNVEGHSVAMFPWYSIP
		l			ĺ	FLNPPCSHTRPSNLPVTQWPPTR
		l			İ	ENNLPSWQLLLTSIHQAQFLSA
1	1	1				LHKEOGSSSEKDGRSPNKWDK
1	1	l				DHIRCPMS\AVMIFSKRHOAL/V
1					i	RAHQGHPNQDNWTVSQMLSK
		1				WWYTLGPNERQKYHELAFQAT
	ŀ	ļ				AALQAHYVHIFPSKGRWELLH
		1			i	GSQRLDWGRCQLKLLGQQGQ
		l			ł	DVLHTNLGKEENSSCWGYKME
		1				CEEGFGPOGGDCGLGGVHAPL
	l					LSPEGLDSEONTLIMGDIFHCA
		1				KKLP
13981	44349	В	14063	1	753	
13982	44350	Ā	14064	2	1457	YPRRRPSRAGVLAGPAVLAGPP
				_		SSPGRRPRRAAFFAEPPFSARRS
	İ	ĺ		i		PRRAAVLAVPPSSPRRRPRRAA
			ŀ		i	VLAAPRVLAAPPSSPRRRPR\PP
			i	1		PSSPRRRPRRAAAGSLPGAELP
		l	l	!		OSSRLAAHL\AAD*PDS*IAAGA
		l		İ		
	ı	ĺ	l		ì	VPRGWEGLTIMAEGKEEQVTF
						YVDGSRQRAYAEKPPVVKTIRS
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP /IPFQFTFQIPSLLQAALWLCL*V
1						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP /IPFQFTFQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP /IPFQFTFQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP /IPFQFTFQIPSLLQAAL WLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP PPPQFTFGUPSLQAAL WLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQ8AGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWPFHPSKQHPRNGLAF
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP IPFQFTFQIPSLLQAAL WLCL*V TWHFFVX*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFGE CPLERVWPFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSGM*/ISSPNSP //PPQFTEQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLFA
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP IPPOFTEDIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQE CPLERVWPFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLPA QHSHAEVLHFGGAMSGQLSLV
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP //PFQFTFQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHICFSPGGLF SISLQSAGCFAGSLQAFEVLP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLFA QHSHAEVLHFGGAMSQQLSLV GPQDSKRTARLTDSQHLGSPT*
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSGM*/SSPNISP //PPQFTEQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PL**APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLPA QHSHAEVLHFGGAMSGQLSLV GPQDSKRTARLTDSQHLGSPT* QPPGLTEGTSLEQG**S*MGWM
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP //PFQFTFQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHICFSPGGLF SISLIQSAGCFAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLFA GHSHAEVLHFGGMMSGQLSLV GPQDSKRTARLTDSQHLGSPT* QPPGLTEGTSLEQG**S*MGWM VGVIAGAPSGWSGSSGPSEGTP
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSGM*/SSPNISP //PPQFTEQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHCFSPGGLF SLSLQSAGCPAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PL**APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLPA QHSHAEVLHFGGAMSGQLSLV GPQDSKRTARLTDSQHLGSPT* QPPGLTEGTSLEQG**S*MGWM
						VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP //PFQFTFQIPSLLQAALWLCL*V TWHFFVR*PVLSSRHICFSPGGLF SISLIQSAGCFAGSLQAEFVALP PGGQSRLHRTRAS*VPGAFQEK CPLERVWFFHPSKQHPRNGLAF PLP*APQRTQPAEDTLPRHPEM GLILSQGTAG/PGRECQATQLFA GHSHAEVLHFGGMMSGQLSLV GPQDSKRTARLTDSQHLGSPT* QPPGLTEGTSLEQG**S*MGWM VGVIAGAPSGWSGSSGPSEGTP
13983 13984	44351 44352	В	14065 14066	72	1235 2715	VVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP //PFQFTFQIPSLLQAALW.CL*V TWHFFVR*PVLSSRHICFSPGGLF SLSLQSAGCPAGSLQAEFVLP PGGQSRLHRTRAS*VPGAFQEK CPLERV WFFHFSKQHFRNGLAF PLP*APQRTQPAEDTLPRHFEM GLILSQGTAG/PGRECQATQLFA QHSHAEVLHFGGAMSQLSLV GPQDSKRTARLTDSQHLGSPT* QPPGLTEGTSLEQG**S*MGWM VGVIAGAPSGWSGSSGPSEGTP LFSSPA*LPWPELFTLC*IGVVR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13985	44353	Α	14067	50	1304	TRKFRRNG*IP*HIHPPKTKSGR
						SRI\FNRPITGSEIEAIINSLPTKKS
				1		PGPDGFTAEFYQRYKEELTNIL
						DEHRCKILNKILANGIQHDIKKL
						IHDDQVGFIPGMQGWFNIRKSI
						NVIQHVNRTRDKNHMIISIDAE
	ŀ					KAFDKIQQPFMLKTLNKLGIDG
						MYLKIIRAMYDKPTANIILNGQ
1	•					RLEAFPLKTGTRQGCPLSPLLFN
1						IVLEVLARAIGQEKEIKGIQLGK
						EEVKLSLSADDMIVYLEKPIISA
						QNLFKLISNFSKLSGYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
	i					ATKRVKYLGIQLTRDVKDLFKE
						NYKPLLNEIKEDTNKWKNIPCS
İ		l				WIGRINIVKMAILPKCKTVGLLS
Ì	l					NISTHKSAVHALVEAGGIPSLIN
ł						LLVCDEPEVHSRCAVILYDIAQ
						CENKDVIAKY
13986	44354	A	14068	1	1155	MFVWRNVEGHSVAVFPWYSIP
l						FLTPPCSHMRPSKLPVTQWPPT
						RENNLPSWQLLLMSVHQAQSL
				i		SALRKEQDSSSEKDGRSPNKW
						DKDHIWWPMS\AVIIFSKQHQA
	İ					L/VRAHQGHPNQDNRTISQMLS
						EQWYTLGPNEMQKYDLAFQN
						VEARIRY/CHRPRKRARFLSHAS
1						PPWRQLPAAGAEASGAWNLPA
1	i					GGG*HDYNRCSVPVFRTEVSTR
						PRPSEICGLTQE*TLLSAKSQVG
1						NTFGFAGHTISDTTTPCGLCTSK AAIEVPEADTLPHICSWHKGEE
		l				CGLRGPEPSSPPPRPSLAARVSA
1						GRGALRRRRRPEVPP\SRAPRSA
						ATATPANEORAVADVGOWAR
l						ARGGRVRAAGRSGLLRVKMM
1	1		l			KMKTVMIHFHLMKDEYIFSS
13987	44355	A	14069	1331	1584	SLQRKAARFASSSTVPRRPQGIL
1378/	44333	ı^	14009	1331	1304	PIISFSNVPVGSRRLKAPLVFIGP
1	l	l	1			G*ITLNRIPYLPHSAARDPFTWN
		l				TFSFLLPCEEVPST
13988	44356	В	14070	ī	1248	II SI BEFCEEVI SI
13989	44357	В	14070	1	459	
13707	44337	டீ	11-071	l'	707	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13990	44358	A	14072		2156	MGKKQSRKTGNSKNQSASPPP KERSSSPAMEQSWMENFDDL REGERGRONYSELKEEVRTHIGK EVKNLEKKLDEWLTRITNAEKS LKDLMELKTTARELHDECTSLK RVSVTEDQMNEMKREEKFREK RIKRNEQSI,QFIWDYVKRPPLH LIGYPESDRENGTKLENTLQJ QENFPLVRQANIQIQEIQRTPQ RYSSRATPRHIVRFTKVEMK EKMLRAAREKGRYTHKGKPR LTADLLAETLQARREWGPIFNI LEKMLPGARESVYAKLSFIDRST RQKVNKDTQELNSALHQADLI DIYRTLHPKSTEYTFFSAPHHTY NCLSDHSAIKLELRIKKLTRNEM AEIKMFFETNENKDKTYQNLW DTFKTVCRGKFIALNAHKRKQE KSIGHTJSQLKGLEKQEGTHS KSIGHTJSQLKGLEKQEGTHS KSIGHTJSQLKGLEKQEGTHS KSIGHTJSQLKGLEKQEGTHS RLINKREENQIDAIKNDKGDIT TDFTEIQTAISEYYKHLYANKL ENLEEMDREDTYTLPRINGE VESLEMRPTGSEIGAINSLPTKK SPGPDGPMRRILPEGGGGGTI PSETVPINKKRNIPPSEIGAINSLPTK SIMLIPKRKGNPPPHFMNSA SIMLIPKPGROTTKKENFRISA SIMLIPKPGROTTKKENFRISA SIMLIPKPGROTTKKENFRISA SIMLIPKPGROTTKKENFRISA
						KAYPP*SSGLHPWDARLVQLTQ

1732

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amine acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13991	44359	A	14073	3	1794	AMLPMELGCGPLPEPLPVGCSR
		1				FSLFK*QTCISTVP/GYMVTAQS
					İ	MSSTPPPPSPSTLPSSPSPPPPLPC
						PLPPPPPSPPTLSSLSSPSPPRPPL
				1	į .	VSPSTLPSPQPSSPQPLLPPSSSPI
		1				SLPSPPPPSPPLPSPSPSAIPSLPPF
				1		SPQPLPPPPPSSPPPSLPSPLLPPP
						PLSSSPSSPLSPSPPPPSPPPSLPPS
				1		PPPSPPPPPPPPQPPSPPSSPLSSPP
				i i		LSSSQPSLLPPSLSSLPLPSSPSPL
				i		LPLSLPLSISPP*LSLLSPLPPSPS
				1		LPPSSFQST*TIGQCFSL/VMWH
		1		1		VAPCTYLALAGNTLMAWPLMS
		1				ASSKASGGVSMFVWRNVEPCS
ł		1		1	l	VAVFSWYSVPFLTPPCSRVRPS
1	1	1				NLPVTQWPPTRAKNLPSRQLLL
				1	l	TSVHQAQSLSALCKEQDSSSEK
		1				DGRSPNKWDKDHIWWPMSGG
				1		HDLQQAAPGPGRAHQGHPYQD
		1				NWTISQILSERWYTLGPNEMQK
		1		1		YHDLAFQHMAGEDIASDEEHM
	l l					VIHEEGVMVSLLMTALAPLTL
					ISSSRIFGKVYGPTPSSSYTYSD	
				1		ASSSTLAPTSFLLGPGAFKAQES
						GEEAEDGLRELETEKALSSSL/R
						RALDQ/*LALIMQLFQAHCFFLS T
13992	44360	Α	14074	1	2496	MLLSIDAEKAFDKIQQPFMLKT
				1		LNKLGHGYEQHHLHTTNDVDE
				1		EDLSDAASKGDDFALSEQSQD
		1		1		AHFLQPEAYGLGEGAETATGT
				1		AHQGNHVRVEECGRSLCGCVP
			İ			LVLHPLPDPSLQPHEAQQPASH
				1		SVACNQRKQPAKLPAVAHERP
						PGGTGSVDPGRPPGATCPESPG
						PATPHTLGVVEPGKSSPPTMEE
				1	1	EPWAPQGSPCWTAQSLSALRK
				1		EQDSSSEKDGRSPNKWDKDHI
						WWPMS\ALMIFSKRHQAL/V
13993	44361	Α	14075	4317	5099	KMFVFLCISAQLSLQCFRGSFHF
		1				IKIYDFSFQLFFLFLELPYLLLLN
						HFKMLELVLQQGHPNQDNWT
						VSQMLSKWWYTLGPNERQKY
		1		1		HELAFQVKVAH\CNKD*KKFSS
	1			1		EAKPTSQGLAG\GNKGSWEWS
		1		1		MSETGTATAPGVSSELLSVAAQ
	1	1				TLQSSDTK\SSFCGAEWGHP*GI
					1	GCDDVIADDGFSTTDTDLKFKE
	1	1		1	1	WVTD\ESGDNSGEEPEGNKGFG
					i	GKVFAPVIPSSFT/HCRPLLDPE/
		1				PPGSPDPPAAFGKVYGPTLSSSY

SEQ ID	ISEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13994	44362	I _A	14076	li li	637	MSRVGSCRWVCGLADFKNGAT
13774	44302	l^	14070	ľ.	051	DLCDKSLHNRELSAERPLNEOI
						AEAEEDKIKKTYPPENKPGQSN
						YSFVDNLNLLKAITEKEKIEKER
						QSIRSSPLDNKLNV/EKDVDFNQ
					1	EFGKLIDGY\DSTKEWDWVHK
				ľ		ODIPNVLVGTAFNVDDGNGGF
					}	VVHWLNNKEFHFTSSTEVFMH
1	l					QLRKLSDKQVDHENDDADRED
1		i i				EEHSQEDRERGLHMKLDH
13995	44363	A	14077	448	464	KKRA*GSVSQRFPAET*LAD*VI
		1				NTGSQEAKRREQR
13996	44364	Α	14078	179	382	NTPNQRSVASNRFVKCTHQHS
	İ	1		ì	İ	VKCTQSALCKMHQSAGFLKVD
		1		1		NHRED*KKGHSDRAKTEHGRG
		ı		ł		QIRE
13997	44365	A	14079	697	1654	PKPSVGRMGFLGTGTW\ILVLV
		1			1	LPIQAFPKPG\GSQ\DKSL\H**ED
		1	j	1	1	LSAERPFELNRLLEAEGRTRFK
	1	1				KTYPP\ENKPG\QSNYSFV\DNL
1		1				NLL\RAITEKEKIEKERQSIRSSP
1		1				LDNKLNVEDVDSTKNRKLIDD
	1	1		1		YDSTKSGLDHKFQENETADNSF
1	1	1				SQEEEPVVAGEDLPPPSPQESD
1	ļ	1		ľ		VQPVQPEEVSARDFLQRLDLQI
		ı				KLSTQAAKKLKEESSSLERMRE
		1				ETSLSPTVPTKRWDTRILLIGAF
	1	1				YRVLIGAFYRVLMGAFYKPIAS
		1				YRALIGVFYRALMGAFYNPLV
		1				RQKSSPRPHSTQEVQLASPLTN TQITYYVIFTV
13998	44366	A	14080	72	1117	ILENHLSDRVAEGGPLOGLRLG
13996	44300	<u> </u> ^	14000	1/2	[''''	VQGEAVPPGLVHQRGGLGPLD
	1	1				GELIGSVGQQVSDVEGAGSHR
	i	1				G*RHTGRTPAVSAGPAA*SATH
	ļ	1				TAPMWLAGW\YEPAQPGPVLV
	1	1				SPPTPFGQTRQASWCSA*ARCA
	l	1				GSWWH*HOTPGGA*RVAAPAP
	į	1				SWSRSCP*RPTRSPAAAPPAPSR
		1				PGRAVGLATGAGHPAOHGHGR
	1	1				AVPTGPRAAGAGSWSELCPPSS
		1				CAGTSRGR*AGTRR*AAPAGSG
		1				RPGPGAWAGPRCSRSTYAAAA
		1	1			CGRGGA*AGADSGPRSGTHAG
	1		1			HCGARCRSSTPLVPAAAGHLPP
	1			1		PGPGRTGDCPASSCPGPGLPHG
	1					CSLSGSWGEGWVSAARQAGLP
	1	1				WAVA
13999	44367	Α	14081	2	269	
14000	44368	Α	14082	1	591	
14001	44369	Α	14083	2	477	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codnn for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
14002	44370	В	14084	237	419	
14003	44371	Α	14085	2	1220	LPPGFVMAAAAARWNHVWVG
					İ	TETGILKGVNLQRKQAANFTA
						GGQPRREEAVSALCWGTGGET
						QMLVGCADRTVKHFSTEDGIF
			1	1		QGSETLPG\GEGMFRGLAQTDG
						TLITCVDSGILRVWHDKDKDTS
						F*PTPGTESGPWGCVGCAKTQA
						HPHVVCHRLGKRMLLKIWDLQ
						GS\GNLCFKAKNVRNDWLDLR
						VPIWDQDIQFLPGSQKLVTCTG
						YHQVRVYDP\ASPQRRPVLETT
						YEEYPLTAMTLTPGGNSVIVGN
						THGQLAEIDLRQGRLLGCLKGL
						AGSVRGVAV\HPSKPLT*PPVRL
						DRVLRIHRIQNPRGLEHKVYLK VSIE/RALLLSGRDNLGG*APEP
						SKTOTRCP*EDTEDR*TLGDPW
						ROLPSGKLFGVWSSPKELSKRD
						GERRSGLGPPAPDAPVPTL
14004	44372	В	14086	96	592	GERRISGEGITATEAT VITE
14005	44373	Ā	14087	102	612	HCAGSPHSPRSLSGTPLPGLPL
11002						WRHLRSPSAHRCTVGAPFWAG
						ODRSOLP*LAGRCGGRGPSGNP
						GCA\PACGPAGVPGGRGLGGPR
		l				TRSSQPALPAPGNEGLSTRASG
		ĺ				CRGRTGSPSSASPPLGLSCLPAG
						QGWGP/GSPPCVSLPPTPWAPV
						RPEPPG*AAPPAPRRPVP
14006	44374	С	14088	1	636	
14007	44375	В	14089	31	612	OF LULY OF OFFICE OF LINE OF OR
14008	44376	A	14090	234	504	SFYHLGAGE*PGKSRVPIPQFQS PWGPYCOGRARGGEVCHLOAV
				1		PPRGAGAARTLSGHPLGSPEQL
1						POPGPAAELPGGVVPRSTPPCP
14009	44377	В	14091	1	924	POPOPARELPOGVVPRSTFFCF
14010	44378	A	14092	11	354	RLSTSPPDSGAQLASPSGSHTRA
14010	44370	l^	14032	l''		AGGAACQSRAVRS/PFPSPWVV
						DGTGCGGAGGGARWGGSGCT
						GAHGVGGR\PGMAG*GPEPCP
						EGROLRPLAPVGCGO*REP\G*K
						NSMGETTT
14011	44379	A	14093	147	513	GSASGVVRLSRWARGLAGFRS
			1			EAADLRGECYSS*GSASGVVRL
				i		SRWARGLAGFRSEAADLRGEC
l				l		YSS*KQRGPKELRSPAGFT*WIP
						/LPGLQVELPASPAPCAPTPQPL
				l	1	SG*WD\GRRGAGGGARRGGSG
1	1			l		RTGALGVG\GGSGMAGCRSRA
l						LPPREGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknowo,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible oueleotide insertion)
				sequence		
		<u> </u>				
14012	44380	A	14094	418	1026	WNPHQGCRWSCLPVPRHALAF
						LSPWVIDGTGRHGAGGGALWG
ĺ			l			GSGRTGAHGVGG\S*GMAGCR
1	1		l			SRALPHGKAAKAR*EIERSASG/
					ł	TGTAGGPSTPSAATGPGAKSLI
					į	ARGQQGWPAAPSAGPAKPTPT
1	ŀ	1				GNSSWPASAAGSSGSCSRLS/SP
		l				HLPAS*GSGLQPWPAQKGAPTV
		1				QQQAEGLPIPRLPEEWISLPFLG
		١				HQGEHSGMA
14013	44381	Α	14095	897	2111	PPLLQSSPNFFLICYLPWHNSHE
	Ì					NTHRTGRRGAGGGARRGGSGR
		1				TGAHGVGGRLRHGGLQVPSPA
		1				PWEGS*GPARNRAQRRYHCSCP
1		1				GLQSGLPHYSGYHT*PDEKEKN
	1	1				GREGQKLER
14014	44382	Α	14096	1304	1922	KLNIGVIYSSEIPDSGAQLASPS
			l			GSGTRAAGGAVCQSCALRSHSS
	1				ALGWSMG/PGRRGAGGGARW	
						GGLGRTGAHGVGGRLRHGGLQ
			İ			VPSPAPREGS*GSVRNPAQRRW
		1				AGTAGGPSTPSAATGPGAKSLI
		1	İ			TRGOOGWPAAPSEGPAKPTPTR
		1	ŀ		}	NSSWPASAARSPGSRSR/RLPPH
		1	ŀ			LPAS*GSGLOPWPAHYFEFFLPI
	1	1			İ	GONAOTEVSY
14015	44383	Α	14097	318	533	MVCFDGADCPWVVDGTGRRG
						AGGGAHRRGSGRTGAREAGGR
			Ì			LRHGGLQVPSPARREGS*GPAS
		i				PAGCLAHL
14016	44384	Α	14098	346	957	ARYTLHIPTRLGSPAGFTQWIPH
						RGCRWSCLPVPRRALAFLSPW
						VVCSFPVDGTGRRGAGGGARR
						GGSGRTGAHGVGGR\PGMAGC
						RSRALPRGKAAKARREIERSAG
						G/TGTAAGPSTPSAAAGPGAKS
l				l		PFARG/LAGPAAPSARPAKPTST
l						RNSSWPASAARSPGSRSWKMS
						LYMKKSASGQSQAQWIAKMW
ŀ						KSPRFKPLWLTS
14017	44385	A	14099	124	663	SCLPVLRRALTFLSPWVVNGTG
14017	44363	Ι^	14099	124	1003	RRGVGGGARRGGSGRTGAHGV
					l	GGR\PGMAGCRSRALPRGKAA
	1		1	l		
l			l		l	KARREIEHSAGG/TGTAGGPGT
1			l	l		PSAATGPGAKSPIAPGQQGWLA
1		1	1	l		ALGINGAGERLFMKVAFSLKTC
		1				SLSCSYLRLTLSRGGSVHVLRG
			1			NLTVCKPQPALLRVVFHHQRQ
		<u> </u>				ADCGALPVP
14018	44386	С	14100	1	672	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
14019	44387	Α	14101	3	1759	RPLRGWRRTRDARAPRRCRCG
						SHGARRCLRKTAATSRCGGAP
			i			GP\RSRPPSTGQAAEAGDW/PAC
		ŀ				CNP*GGLRAADGPGSRPPRGSI
						GLLCGQGGQGGSSPWVVHGTG
					1	RCGAGGGAPWGGSGRTGAHG
						VGGRLRHGGLQVPSPAPQEGS*
					i	GSARNRAQPRWAGTAGGPSTP
						SAATGPGRDGEGG\ETGAHVSA
			ĺ			LKGTPGTEWPAHGLVPYSIGDQ
		l				TCLIFKVTFKAVVLKCGPQTTS
		l				STSISCPGSAN/SPPSPAG*/GSSG
						CGSSSLCSAHSVQEDSQPQ*KR
		1			i	TGLQS\TYGRSGSSSNEPRYHSD
		1				LPTSRSKLRNEDTFFGRRKKLS
		1			1	EVQRTGQDPA*SHHTSSYTLPG
					1	L*TAEGKDRVENAA*KGSPNFQ
						RSGTFNPKGGARKTRQAWHPR
		l				APNCQAGTPPREGSSGPRTSLS
i		1				AAFRIPQEVVNFLISGPITRAQR
					ŀ	HGEPESPRGQTGSPPLGKTGSE
		l				GTDRKSTGADKKSTGDRQEVR
						EPTESPQGLDRKPTGMDRRSTG
		1			ŀ	NRQKVRRQIGSPLGLNRKPNGT
1						DKKSTRNRQEVRRRQSAGARQ
1						ERAGSSTFLRLSALCGLLVDVN
						REQVAKQKWDLYSSRFTSGNR
14020	44388	A	14102	1879	2607	GKPSLSLHPGHKCGISRCNGNV
		1				SSRLRSPAGFTQWIPHRGCRWS
		1				CLPAPRCALALLSPWVVDGTG
		ı			1	RHGAGGGARRGGSGRTGAHG
		1				VGGRLRHGGLQVPSPAPREGS*
						GPARNRAQRRWAGTAGGLSTP
						SAATGPGAKSPIARGQQGWPA
		ı				APS\GGPPSPSPPGTPAGPQAPH
		1				AAPVPAR\PLPPHLPAS*GSGLQ
		1				PWPAQKGAPTVQGGG*RAPQM
		L			l	PPKWEPRQGRRIVAFTIEVGVL
14021	44389	Α	14103	2	272	GHWGMVNSPCRASATVSRGSP
						LPGVLPTSPPSATSTCRMRCHPR
1		1	1		l	RCPLPFRRLSPPSLTPP/PGKPPL
		1	1			LPRPHSMPGLGLPLGGLPVFHP

sequence 09/540,217 c	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
s		of peptide sequence	deletion \mpossible nucleotide incontion)
	sequence		ocietion,possible nucleonide insertion)
14022 44390 A 14104			
1022 11350	101	1815	OSAROLRPCPGAPSLAGVENRE
			KKTTNLTSLAAPDTQSTCGQSC
			GCLHSMGSSCMQHSYYTGFLR
			SLWPCRCHRSAFSGQRRSEEKD
1 1 1 1			SKSVRERSGRSQNVPARPALTS
			PLTQSPGPLAPGAGSWSRSQLG
			TGTGVLWWQRESATGDFRTES
		1	ARSSDPSRRDWPGAKRKREEG
			AATVLWACATASTAIICODNAI
			SPLKSGNSPSNHLWRPQGQLPP
			VLGLAQPKRWRPNRLTTSPSPS
			RMMARPVDPQRSPDPTFRSSTR
			HSGKLEPMEATAHLLRKQCPSF
			LNSPAWEASGLHWSSLDSPVG\
1			PCRP*GLRPSTHGAR\TFSGARP
			GLGGYSPPPEEAMPFEFDQPAQ
			RGCSQLLLQVPDLAPGGPGAA
1			GVPGAPPEEPQALRPAKAGSRG
!			GYSPPPEETMPFELDGEGFGDD
			SPPPGLSRVIAQVDGSSQFAAV
1 1 1 1			AASSHQCLSHHIPMSPSVGRNG
1 1 1 1			LLTKPPSTSRRGKLTKRDDMYY
1 1 1 1			FVPFRSLNDMMHMLPFIEDSR
1 1 1 1			WTHNLDINREQKTNPLAGPKG
1 1 1 1			RPISRTQAAASTSNSKTESQIQE
			KTEEAPFSLRYININNRSMDPLN
14023 44391 A 14105 3	39	877	AGPGPPRPLPPRSSEGPPGASISL
			CSEVPGPSCQGRFRGGRDAMS
1 1 1 1			MPWREEERSLLAGE*PS*GG*S
1 1 1 1			GRGVSRKCS*PGPRGVPRPRPL
1 1 1 1			RSRPRAASSE\DQNSPLDQSLHR
1 1 1 1			TRKTIPSSGGSTAIMPRRGSSSE
1			TFFSAPSKTWVLGIPGETFISGP
1			AAFP\APGGPRAAPRPSPSLNSL
1 1 1			*SGGNGGPGPPRLHFSSCGKGP
			PRPSNSRN\GGQ\$ACQGGSHLV
			C*SHRGSLGEGHAYQGDPCSPG
1 1 1 1			HPAKAPSRAHAQARRSAHAAS
			LTSGPCDASCQAW
14024 44392 A 14106 3	358	546	CFPSRIRLALSRAPKWPAKRSPS
			EPEMS/PPVWSPHPDPNNTAPPK
			EQPPSSPGRVGRHGNWSV
14025 44393 A 14107 2	22	549	DHCLCSVVRNKRDSVYPLPIPP
			CCKTSTISPP/LVDAVRP*DISVS
			KRVTSCAPLEPQPHPCLPTINAG
			R/TLARPQGAPTPNKEEGQAPG
1 1 1 1			RATASGQAAPATTVLGDTTK/G
1 1			PQGRHRGTSASAAGPAPFAVA
1 1 1 1			DQLPPEDRAASAEGRK*AAAFP
1 1			FCHRLVQDGGPPPPAAGVPGA
1 1 1 1			GES

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
			1	sequence		
14026	44394	Α	14108	146	2616	LAAVGMASHCFRQRRQSQSPA\
		l				GLRGAGSAYSPA\EGARSSHGQ/
		1				PPWRGRRRGPGDPVRGAARCR
		1				RPRAAAQPARPAARPPARPRPA
		1				P*AACCE/VPSAANSVAPSAAM
		1				*SSAGRDGASSQRSRRPHAAPP
						RSSALPANGK/PPSPVTRAARTC
		l				GARSSNTGA*SAARPKHSAASS
		1				RHPPPAA/MPLPARHA/PGGTSG
		1				RPLPTPQGSAPTSSASPRGPATP
		l				PPAGPRAPPPSPRRAAAPRPPGR
		1	İ			QLSPAPGTPAAGGGGPPFPALR
		l		1		*SGSVFGRQLVGDGERGRACG
		l				ARGSAAVPALRTFVVSPSTVVA
i		l				GAACPLAVALPAPDPLPCSE*A
		1				PLRSG*VPQPASLFCSQFRPQA
1		1				WQHP*LLRAPCPRLGAAVPCV
		1			1	WCHSRALDSATPRWM*G*RLP
		1				AVWL\PAFPSPPFNQAFVPPCQQ
		1				NCSLQNRQTLSASAKIGTEEWD
		1				IAIKIPENVEVTLELGNEQRLEE
		1				FGGLRRRQEDEGKFGTPTDWL
		1				NGCDQNADSNMDSEGQGEDG
			i			GCSFGGAGLRGSGWGDQTRET
						FLALKANLRWPFRSSAQSQTSQ
		l				EGAAGARSGLKGGKKNPLKLP
]				KKQAKKMNKAFKQKQEKERK
		1				KHKELKAKASWKGLRPQVEFK
		ł		1		NLAERRGEAAVPGCPVVAGHS
1		1				WRLLSLARGCAKRARQLLFPSE
		1				KLLAALREGSTRGGGAASQPQ
					004	RKLRVCMRVLDSSVEDFHMDG
14027	44395	A	14109	1	885	MIISIDTEKALDKILHPFILKTLN
	1	l		İ		KLGIDGTYLKIIRALYDKPTANI
		l				IMNEQKLEAFPLKSNTKQGCPF
l		l				LPLLFNIVLEVLARAIKQEKEIK
1		l				SIQIGREEVKLSLFADDMIVYLE
		l				NPIISAPNLLKLISNFSKVSGYKI
1		l				NVQKSQAFLYTSNRQTEIQMM
	1	1	1	l		SELPFTIATKGIKYLGI\HPLTRD
		1	1			VKDLFKENSKPLL/KELEKTTLN
			1			FIWNQKRAHIAKTMLSKKNKA
		I				GGVMLPDFKLYYKAAVTKTA
		l				WLGDDQVFLIPRGHISDYHMG
		l				KNLGQYPAFQDRGPCGFPQCIV
		<u> </u>				PLVYRD

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide inscrtion)
		l		sequence		
11000	Lucas	 	11110	Lisa	891	L LIHLSTDALRVSGWGTVRSFSS
14028	44396	Α	14110	453	891	
		l l				HEAIFQTIAWGETLDNTLLSRA
		1				EVPAAFHGALCPWFIETREWQ*
		I				LLQSILLVNIWLTRHVLHSPRSL
		l				KP*FYTT*VFVCSKLGQSGWGK
						VAGAKLQMNNISAKQLFKVQV
		ļ.				LFKIESLMSSLST
14029	44397	Α	14111	452	2922	KMGLAREWILRFLVLILQEIRS
ł		l				MRQKENKDIQDLNSALHQADL
		ŀ				IDIYRTLHPKSTEYTFFSAPHCI/
		ŀ				RTYSKIDHTLGSKALLSKCKITE
l						IITD\$QTTVLSELRIKKLTQ\NRS
l		ŀ				ATWKLNS/YLNDYWVHNEMK
	Į.	l				AEIKMFFETNENKDTTYQNLW
		i i				DTFKAVCRGKFIALNAHKRKQ
	1					ERSKIDTLMSQLKELEKQEQTH
1	İ	i i				SKVSRRPITG\SEIEAIINSFTNPK
l						RVPGPDGFDEPNFYQRYKQEL
1	1					VPFLLKLFQSIEKDGILPN*FYE
						ASIILIPKPGRGTTKK/EENFRPIS
	1	ŀ				LMNIDAKILNKILANRIQQHIKK
İ					1	LLHHDQVGFIPGMQGWFNICKS
					1	INVIQHINRTSDKNHTIISIDAEK
	1	1	i			AFNKIQQLFMLKTLNKLGTNG
	1					MYLKIVRAIYDKPTANIILNGQ
	1		Ì			KLEAFPLKTGTRQGCPLSPLLF
	1					NIVLEVLARAIRQEKEIQGIQLG
				l		KEEVKLSLFADDMIVYLENPIV
	l .					SAQNLLKLISNFSKVSGYKINLQ
	l .					KSQAFLYTNNRQTESQIMSELP
	1					FTTASKRIKYLGIQLTRDVKDLF
	1					KENYKPLLNEIKEDTNKWRNIP
						CSWVGRINIVKMAILPKVTYRF
	1					NAIPIKLPMTFFTELEKTILKFR
1						WNQKRAHIAKTILSQKNKAGGI
						RLPDFKLFYKATVTKTARYWY
						ENRDIDQWNRTEPLEIMPHIYN
						HLIFDKPDKNKOWGKDSLFNK
						WCWENWLAICRKLKLDPFLTS
14030	44398	A	14112	3	3349	
. 4050	1.1370		1 - / - /	17	I	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14001	44399	<u> </u>	14113	294	1107	RRHRLENYIPRTAOSPHPRKAL
14031	44399	A	14113	294	1107	
						ARSESKRDGGFKNNWSFDHEE
						ESEGDTDKEPGPLDAPPRP/ESG
						PRAPGKGL*RRSRSPGPSERGSA
				·		ARLTAEPEARNPAASTLLRRGP
						SSRRARRRCGPWIKKEAAWEE
						EDGGRQERRLGQDLRTRTLNK
						SKGGIDSRIISLGQHSHPTHAKL
	ļ					WLDQSLREMEVLKIIGALIMKK
	ļ					KVKEIQIKSRARSTPLHDQIGTT
	İ					SPRERTIASKRHPYYITPKGIRY
						EQDTLSRTPITTIRSRQRPLGPLT
						ASPPPSFPTLAPPTQAIKPSLHLR
	ļ					EGKPTPTPGERSTRPHHPEPPRK
			1			EYALSSLYPPTYLLAPTSTDLG
			1			NAHPTQEPQFNVDATHPRQNR
						FIPVAGNSPKRSSQFPIVFAPKK
						NTTPPYLDFAAILGLHRRPPRSR
						FRTSYPPKEFRSSPPERTAPRN
14032	44400	Α	14114	2	297	VQKSVLCSFQAGTMEGVEEKK
		1				VPAVPETLKKKRRNFADAIQFL
			l			QTVFLAF*YTLLKGRQ\VYWLV
			i			EE*HQFYKISYRKKMVQRIRQH
	İ		İ			PQSCTVQKSWWP
14033	44401	Ä	14115	3	754	GTMEGVEEKKKEVPAVPETLK
				i		KKRRNFAELKIKRLRKKFAQK
			İ			MLRKARRKLIYEKAKHYHKEY
						ROMYRTEIRMARMARKAGNF
			İ			YVPAEPKLAFVIRIRGINGVSPK
						VRKVLOLLRLROIFN\GTF\VKL
	İ					NKASINMLRIVEPYIA*EYPNLK
						SVNELIYKHGYGKINKK*VALT
1				l		DNTLIARPLGKYGITCMEDL/IH
1						EIYTVGKCFKEANNFLWPFKLS
						SALSGMK/KKTTHFVEGGDGG
		l				NRVDQINTLNRRMN
14034	44402	A	14116	1	388	THE POSTATE TRANSPORT
14034	74402	1~_	14110	1	300	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ı		sequence	1	
		<u> </u>				
14035	44403	A	14117	1	1145	MTRNYEWLLEAESSPQPIAFQK
						MEPQPYNSKELNSANNQCLLE
						KPKLRQRLKDKEYDQGHTAVQ
			ł			CQSWVYVLTPDSFCQQLILLPS
						TENFHECSRSDTAGHLRTGHVA
					1	SNWLSEHSPQTMPAAMQWDK
						KMQNTSSWFPQKAPEKVKPMS
		1				KLWQTPSQNDTLYKAVKALTL
				1	1	SSPERNSINTNKKDVHTETSSK
		l				GHQHQRPKVDKSMKMRKNQH
				1	ł	KKAENSQNQNASSPPRNHNSSP
						ARERNWSENEFDELTEVVFRR
		1				WVITNSSELKEHVLTQCKEAKN
						LEKRLEELLTRITGLEINQAEERI
		l				SETEDQLNEIKHEDKIREKRMK
						RNEQILQEIWAYVK\RPKLRLIG
1						VPENDRENGNKLENTLQDTIQE
i						NFPELAR*ANIQIQ
14036	44404	С	14118	291	551	
14037	44405	В	14119	1	1308	
14038	44406	Α	14120	82	367	ITLSSPPHSFIGHKTPASGPPQKI
	1		1			PPQPSQ/CKQWGDPTPVEAC/CV
	1					SCPTQGFISPTRGGPPCKHESLV
	1	ĺ				VPLVGTPRRNQPGGHIQHSQR
					_	WAPVLT
14039	44407	Α	14121	999	2661	VHLTGCGGPGQPVPEPAPPRGL
1	İ	ľ				RSMRC/GP*GFSPGAEA*TEP/C
						HYGGRGFPCGPFSLGCLAT\NSP
	ł	ļ				WLPGPRPPCSPWAGP\CGLCPLP
ł	l	l		ĺ		AAAAERQPALSVPGASPSCRLL
		ŀ				PLVVLCHGSHLNGGRVLLRPG
						GSPCGHCHNCAGARRGLGHVL
		L				QGGAQARQREQLQR
14040	44408	В	14122	164	1042	
14041	44409	В_	14123	320	566	
14042	44410	В	14124	1539	1674	
14043	44411	Α	14125	185	780	QKAKKGKLKRSKQNSFPGLKK
1	1	ŀ				TVPSQLVQYCPSF*ISWKPMVY
		ŀ	l	ł	1	QKVPA*PRPPRTPDHHPDPGPA
1		ĺ		1		PHPRQHHPPPRSPDAEP/SPTTPT
		l				RRRPPTQPPDPHPKRPPPHSKPH
		l		l		PPQQPPTPPTATPTEQQRHTAPP
						TPRPPTTHPPSTPKHPA\PHTAPP
		1		l		PTKNTKPPTPQRTTQDKGRQKK
		L				PRRQPHHSEPNKKSPKAHI
14044	44412	C	14126	1	1128	
14045	44413	Α	14127	319	536	
14046	44414	С	14128	231	425	
14047	44415	В	14129	1	1431	
14048	44416	Α	14130	75	393	
14049	44417	C_	14131	174	371	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codnn for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14050	44418	Α	14132	79	164	
14051	44419	Α	14133	343	775	
14052	44420	Α	14134	554	724	SEGPSKEPTHRRMKFSHHDDLI
	ì					FLDLSN*QPPNSPEPHPPQSP*KP
		l				WPQNPSSRQM
14053	44421	Α	14135	20	242	VSFLSMGSGHCIRSTRGSKMVS
	1				1	WSVIAKIQEI*CEEDERKMARES
	1					LAESMSTYVMMNHTYDS*GK
	1	l				THILIMKK
14054	44422	Α	14136	229	366	
14055	44423	Α	14137	131	505	
14056	44424	Α	14138		1317	AGCSCGSRAMAEEQGRERDSV
						PKPSVLFLHPDLGVGGAERLVL
	i					DAALALQARGCSVKIWTAHYD
		l				PGHCFAESRELPVRCAGDWLPR
	l .					GLGWGGRGAAV\CAYVRMVFL
		1				ALYVLFLADEEFDVVVCDQVS
	1					ACIPVFRLARRRKKILF\YCHFA
	1					DLLLTKRDSFLKRLYRAPIDWI
	1	1				EEYTTGMADCILVNSQFTAAVF
	I					KETFKSLSH\KDPDVLYP\SLNV
i	1	l				TSFDFSCS*KAGMT*SPRGKNS
		1				WLLSINRY\EREGKIWTLGTGK
	1	l				ALVQLRGRLTSQDWERVHLIV
		l				AGGYDERVLENVEHYQELKK
		l				MVQQSDLGQYVTFLRSFSDKQ
		l				KISLL/RQLARVCFYTPRQ*GTL
		l				GIVPLGRPCYMAVPQFICCLIRV
		l				GPFGSSI*PQCSQGFCVEPDPVH
		1				FSEAIEKFIREPSLKATMGLGWE
		1				EPE*REKFSPGSILQEQLYRYVY
14057	44425	Α	14139	1	648	MESKEEQGGVADHQKATRGIP
		l		l	1	TPQPREVVSERATQRGNRAFST
	1	l			1	KLCNPKTNRKQQQQYQQQK
						GRTKTLPKGQHPQRSKLDKLTE
						MRKHQQKNTKNPKGQSPCSPP
	l .	ł				NDCNVSPARVHNWTGDEMEEL
		1		ŀ		TEVGFKKWVRKNCAELKEHVL
		1				TQCKEAKNLDKRLEELLTRKTS
	1	1		1		LERNINDLMELKT/RA*ELCKA
	1	ı	l			YTSINS*IDOAEERISEFEDHLAE

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14058	44426	A	14140		1365	MPWKGGTKCIKYLLFGANFIW SERDCRALSØFTCSRAPDTG RGRECRQLALPAPGRHRAGRES ALRLPAGSKAGSRAGBORDG RIRVGVSGTLFGSTTGLFNQSL AETGFTPYTQKSNDFLVKSLKI WHTYFKNFHQEARLYIRTSPGS WNWWFCTLGIESPEQPGYVEG QRPRVVTKSFASLTLKRYGTVL LLAARINADVLYVLQLAGIAVL AIGLWLRFDSQTKSIFEQETNN NNSSPYTGVYILIGAGIAVL AIGLWLRFDSQTKSIFEQETNN VNSSPYTGVYILIGAGIAVL GEIGCCGAVCSSQCMLGLFG FLLVIFAIEIAAAIWGYSHKDEV IKEVQEFYKDTTYNKLTKDEPG LLVIFAIEIAAAIWGYSHKDEV IKEVQEFYKDTYNKLKTKDEPG ETLKAHIPYAVCKLGKDTLLRF ENTLKAHIPYAVCKLGKDTLLRF LRIVSAHRLNCGGLAGGVEGFI LGVEFDKTFTVKSCPDAIK EVFDNKFHIIGAVGIGIAVVMPS FRSNIPPDHVSSYLLUFGMIFS MILCCAIRRNREMY
14059	44427	В	14141	98	1064	
14060	44428	A	14142	560	964	KRLPRILGIRGKPCCPDTSRAGR RVRGRAAAPCREAARGGQRR FLPPTWRCETGAATMFBSPALT PTPFSVKDILNITVRPRS/LGQAA LTKRLSASIJSPQSCARCRRWS CVILSAVRGN*YAGSRAHRDCT Q
14061	44429	c	14143	189	452	
14062	44430	Α	14144	1	647	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
14063	44431	Α	14145	890	2905	LRLSLKFPMSQWTPEYNELYTL
						KVDMKSEIPSDAPKTOESLKGI
						LLHPEPIGAAKSFPAGVEMINSK
						VGNEFSHLCDDSQKQEKEMNG
						NOOEOEKSLVVRKKRKSOOAG
						PSYVONCVKENQGILGLRQHL
						GTPSDEDNDSSFSDCLSSPSSSL
					i	HFGDSDTVTSDEDKEVSVRHSQ
					ł	TILNAKSRSHSARSHKWPRTET
ľ						ESVSGLLMKRPCLHGSSLRRLP
			ł			CRKRFVKNNSSQRTQKQKERIL
						MORKKREVLARRKYALLPSSSS
						SSENDLSSESSSSSSTEGEEDLF
						VSASENHQNNPAVPSVTYGQP
		l			İ	WSKNIKEKTLEINSSOALTAYE
1						CLHSAHVINLISPFYHHFPSSOE
						RSYLSTLSTLSNKKTGSVWFTR
		ı	ŀ			SPKVTOFFPGGAITGTLSORPER
						EILRPLALPGKGNTPPYFCSPSM
	l					GCTHCPTSPNEMNQGISEDTAA
	İ			}	1	APVRDIYIKPPSPWDRAPGGRG
			l			GCGHSFSRLKSPYLRALKSAAD
						FPAQHSSSDKGQAASSSGSLTP
						VYPVWVTPPSRGRQTPHRGEL
			l			WLASDGCPSGRKLPEEGTGSNL
						CCSAASAGDTQANRDHNSSPA
		1				REQTWTENKFDKLTEAGLRRW
						VINSSELKEPVLTQCKEAKNLE
	ļ					KRLDESPTRITSSEKNINDLMEL
		1		_		KNTAQELHEEHTSINSQ/DQAE*
14064	44432	Α	14146	1	737	LNRGEQRAVRYYSHMKLNMA
						EEEDYMSGFLH*CPRRYQTRIA
						NAKANPRSPSKR\EKQQEANLK
						NRQKSLKEEEQERRDIGLKNAL
			ŀ			GCENKGFALLQKMGYKSGQAL
	i					GKSGKSGIGHEASLKRKAEEKL
			1			ESYRKKIHMKNQAEEKAAEQF
						RMRLKNKQDEMKLEGDLRRSQ
			1		l	RACQQLDVQKNIQVPREAWY
						WLRLEEETEEDEEEKEQDEDEY
						KSEDLSIKKTYLQIVQDQLLQI
						MTKIIPNKVET
14065	44433	A	14147	1	1860	
14066	44434	A	14148	1	1203	
14067	44435	В	14149	1	1119	
14068	44436	Α	14150	1	1407	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14069	44437	A	14151		1731	ERSSPATEQSWMENDFDELRE EGFRENFELLEKWTHHKQD KNLEKKSWFFEKINKIDRPLAR LIKKKREKSQINAIKNDKGDITA LIKKKREKSQINAIKNDKGDITA DPTEIQTITIETYKHLYANKLE NLEAMDKFLDTYTLPRINOEEV ESLNIKLINNELN*GINN**LTNY KNYQDQMD*QLNSTRGTRRS WYRFLLKLPQSIEKERILPNSTY EGILLIPROGENDITKKDHSFRIS LMNMDAKILNKLLANRIQOHIK KLIHIDQVGFIPGMQGWFNICK SINVIQHINRTKDKNHMIISTDA GKAFDKIQOFPMLKTLNKLGIDI GMNLKIIRATUNKTTANKLGIDI GMNLKIIRATUNKTTANKLGIDI GMNLKIIRATUNKTYTANKLGIDI KLEAPPLKTGTRQGCPLSPLLF NIVLEYLARAIRQEKEIKGIQLG KEEVKLSLFADDMIVYLENPIV SAQNLLKLISNFSKYSOYKISVQ KSQAFLYTNNRQTESQIMSEIPP PIASKRIKYLGIQLTBDVKDLFK ENCKPLLEKENTNKWKIPC SWIERINIVKMAILPKVIYSFNAI PIKHEMTEFTELEKTTLKFIWNG SWIERINIVKMAILPKVIYSFNAI PIKHEMTEFTELEKTTLKFIWNG KRARITYSIEROKNARGGITLPE
						FKLYYKTTVTKIAWYG
14070	44438	В	14152	1	906	
14071	44439	A	14153		553	EHSSPATEQSWMENDFDELRE EGFRENNYSELKEEVRTHIGKEV KNLEKKLDEWLTRITNAEKSLK DLMELKTTAREICDECTSLSSR CDQLPRPDGFTAEFYERYKEEM VPFLLKLFQSIEKE/VNPP*LTL* GQHHPDTKAWQRHNKKREF*T NIPDEH*CKNPQ*NTGKPNPAA HQKAYPP
14072	44440	A	14154		909	MKA EIKMFFETNENKDTTYQN LWOTFKA VCRGKFIAVNYHKR KQERSKIDTLTSQLKELEKQEQ THISKASRRQEITKIRAELKEIET ONTLOKINASS SWFFETNICH RPLARLIKKKREKNQIDAIKND KGDITTDPTEIQTTIREYYKHLY NQEEVESLNRPITGAEIEAIINSL PTKKSPQDGFTAEFYQRYKEE LECSSSPAMEQSWTENDFDDLR EEGFRSNIFSELKEEVQTHIKE AKNLEKLEKWLTNITIVEKS LNYLRELKTMARRELC
14073	4444I	A	14155	1	969	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
14074	44442	A	14156	377	687	LLEGKLTNRKDIHTKTPSV/SPP
						SSKTKEHSSSPAMEQSWMEND
						FHELREEGFRRSNFSKLKEEVR
				l		THRKEVENLEKRLDEWLTRITK
						VEKALNDLMELKTMARE
14075	44443	Α	14157	1	711	
14076	44444	Α	14158	1	831	
14077	44445	A	14159	1	744	
14078	44446	В	14160	1	1479	
14079	44447	A	14161	1	810	
14080	44448	Α	14162	3	651	_
14081	44449	A	14163	2	1067	
14082	44450	Α	14164	2	2063	
14083	44451	Α	14165	1328	1639	PTNAHETSGRYIAGSSEKRTSA
						RTCGGTSAAARPAGTSMQTRC
		l				RQPPPSS*HAVPRVAQSSDSGL
		l				KATHPQAYQPCGHTPTRLPQES
	1					NCIEHERGLGKKKKKKK
14084	44452	Α	14166	1	620	MTLGYGQRGANKDLRTGFDPD
		1				LCTMDNFAEGDFTVADYALLE
i		1			·	DCPHVDDCVFAAEFMSNDYVR
		1				VTQLYCDGVNDSFLIGGLLRIG
		i				CKIENERSSSPAMEQSWTENDF
		1				DELREEGFRRSNYSELKEEVR\T
		1				NGKEVRNFEKKLDEWITRITNA
1	l	1				EKSLKDLMELKTTARELCDKC
	l	1				TNLSNRCDQLEERVSAMEDEM
1		1				NEMKHEDKFREKE
14085	44453	Α	14167	1	514	MGKKQNRKTENSKNQRTSSSP
						KERSSSPAAEQSWMENDFDEL
		l				REEGFRRSNFSELKEEVRTHGK
		l				EVKNLEKRLDKWLTRITNTQKS
		1				LKDLMELKTTARELHDECTSLT
						NQFDQLEERINNI\EDH*QD**R
1						RKERRIK*MQ*KMIKGISPLIPQ
						KYKLPSENTINTSMQIN
14086	44454	Α	14168	1	366	MDTFLQAERKDYMEAYELIEQ
		l				EEQGEREPVAVNNILSTEALMA
İ		1		i		NAKSTLMFYAVGFLWQPGGGR
l		ı	1	l		GHKAASNPGVTRDVVGSLSSLT
	1	1	l			FLRGERSSSPAMEQSWTDNDFD
				i		EL\REEGFR*SNFS
14087	44455	A	14169	1	1026	
14088	44456	Α	14170	1	3906	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14089	44457	Α	14171	1643	2721	IKPTKMGKKQSRKTGNSEKQST
		1				SPPPKERSSSPAMEQSWMENDF
		1				DELTGAGFRRLVITDFSELKED
		1				VOTHHKEAKNLEKRLDKCRTN
		1			1	G*LEWMNG/ITRITNAEKSLKDL
		1				MELKTKARELHDECTSLRSRFD
		1			i	QLEERVSVMEDQKK\MNEMKR
		1				EGKFREKRVKRNEQSLQEIWD
		1				YVKRPNLRLIGVPESDGENGTK
		1	ļ.			LENTLQDIIQENFPNLARQANV
		1	-			QIQEIQRTPQRYSSRRATPRHIIV
		ł			1	RFTKVEMKEKMLRAAREKGQ
		ł				VTLKGKHIRLTVDLSAETLQAR
		1				REWGPIFNILKEKNFQPRISYPG
		1				KLSFISEGEIKYFTDKQMLRDFV
		1			i e	TTRPALKELLKEVLNMERNNR
		1				YQPLQNHAKM
14090	44458	Α	14172	351	898	SGVTAPWRPNGTLLITAATASH
		1				PVLAIPAHLCSPRCPSFLTTVTC
		1				SGPQHHAARGRVPFHC/SECGK
		1				SFRYRSDLRRHF/ARHTALKPH
		1	İ			ACPRCG/KGFKHSFNLANHLRS
		1	İ			HTG/ERPYRCSACPKGFRDST/G
		1		ľ		LLHHQVKTITT*PD/CGGCLVSN
		1				IVGSYSRPPWPCTRGPRRPLGSP
		1				SPAPGSVCLC
1409 I	44459	Α	14173	1	551	
14092	44460	A	14174	3031	3207	TIHGCWWDPLSSIF*PFWGRGFI
		1	ļ			SHKICQAQISFRVISGTVLSLDK
		l				LLIMFHKAIFASP
14093	44461	Α	14175	3555	3957 .	ENGQKTWTGTSQRRYTGDRM
		1				AQIQNPDTKCW*ECKATGTRIH
		1				CWWECKMVQPLWKS\WQFLT
		1	1			KLSILLPYNLAIILLGNYPNALK
		1				MYAHTKGFTCMFIAASFIIAKT
						WKQPRRPSVGEW/INNL*LVQT
		<u> </u>				MEYYSAL
14094	44462	A	14176	3	883	CQPYPRPRPRTGTHGSCQPPSQ
						GSAEP*G*CLPTLLRMALKAQP
						TKVNPRTLLDSDPDVC/S*AAHS
						RVLITSPSRAPWRLPSP*RRCPA
	ŀ					GTCMPSTATSAR*CSQRPGSSW
	1	1				*T*TGSSLGSQACGPSSSTAPPSS
	1					SWSACTCGCAAAARCSCAASS
		1				TRSGPTCGSSPPASSCASSTPAP
						GTTPSSTLTSWASSPWSTPCPGS
		1	1			AGPSSWSSSSSATPSASASTRTL
	1	1		1		SPGSPAAP*PWPTALSRLTEQEA
	l	1	Ì	1		VWGAWGISHGNSWDKETKLV
		1				RLPHPCSKSPCFRPSLQPYRAHR
	ı	1	1	I	1	AP

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14095	44463	A	14177	3	395	SAEVGAAETTLTELRRTVQSLEI DLDSMRNLKASLENSL\GILLHL ESELAQTRAEGQRQAQEYEAL LNIKVKLEAEIATYRRLLEDGE DFNLGDALDSSNSMQTIQKTTT
14096	44464	A	14178	2	1044	RRIVDGKVVSETNDTKVLRH AKTSSQMPSPQEGMCGKACTP ALSQADSLCPLLRLASEVEGYS LPACAEPYVQSECLSHLSVWSL
						QHALSSLASMSFTTCSAFTNYW SPGSVQVPSYGTQPVSHAASVY AGLGGSGSRISVSHFTNG\GGLA
						GMGGIQNEKETMQSLRDRLAS YLDRVRGLETENWKLESKIQEH LNVTRL*LETEIEALKKELLFM
						KKNNEEEAKGLQAQIASSGLTV EVDAPKS/QDLAKNMADSRAQ YDKLARKNREELDKYWSQQIE
						ESTRVVTTQSAKVGAAEMTLT ELRHRVQSLEIDLDSTRNLKAS LENSLREVEARYALQMEPSLLF
			<u> </u>			SKIIFGRVWEISYLHVFEGKQV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	l			sequence		
14097	44465	A	14179	li	2040	MPVPOLPPPVSRLAATAAASVT
						EPGPNPEAVRSASAEPEVKEGF
		i				PSONHSVGHSNLIPSGGSVAPSS
						GRSHYGYASEDGTCSKLRVRPR
		l				LAAAPAAPFAFVADLASCCQG
	İ		ŀ			LSFFEGILLVRGPYRRLRATVV
	l				İ	RDLVAVRMAEEQEFTQLCKLP
						AQPSHPHCVNNTYRSAQHSQA
						LLRGLLALRDSGILFDVVLVVE
	i					GRHIEAHRILLAASCDYFSMSFT
						TRSTFSTNYWSLGSVŁATTYGA
						RLVSSTARVYAGAGGSGSRISV
						SRSTSFQGGLESGGLAAGMAG
						GLAGMGGIQNEKETMQSLNDR
l					i	LASYLDRVRSLETENRRLESKIR
						EHLEKKGPQVRDWSHYFKIIED
						LRAQIFANTVDNARIVLQIDNA
ļ			ŀ			RLAADDFRVKYETELAMRQSV
l			1			ENDIHGLRKVIDDTNITRLQLET
			l			EIEALK*ELLFMKKTPEEEVKG
l						LQAQIASSG\LTVEV\DAPKSQD
	l					LAKIMADIR\AQYDELARK\NRE
		1				ELDKYWSQQIEESTTVVTTQSA
		1				EVGAAET\TLTELRRTVQVFGD
						STWTSMRNLKA\NLENSL\GEV
						EARYAL\QMEQL\NGILLHLESQ
						LGQTPRTEAQRQAQE\YEALL\N
				1		IKVKL\EAEICHLTRPPSWKIGE
l				1	1	DFNL\GDS\LDERNSMQTIQKTT
		l		1	i	TRRISWIGQSGVLRPIDTKVLEA
		L				LSQQEASGTLLGKQEANKKFQS
14098	44466	Α	14180	275	550	
14099	44467	В	14181	729	909	

SEQ ID NO:	SEQ ID NO: of peptide sequence		in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion)
14100	44468	A	14182		1415	MGTQCDILAKSLEDCPVJEVE DAPQGLGKEDEIEQFSDKTFG SGAIDDWREAHECLAELEVK QPVAVIEQTGNGERDEMDLLD DHEENLAERLSKMYLENELEDP AIMRAVQTRPVLQPQRGSLNSSI, WDGSVYSSEANPRITACSGNAY SVCIRICFASEASPRYQKMIGTF LINEHYQGQULHSLAVLLLELS. P*APHLINRWLYPASPNRFCVRS PNOLCSYPNSSILGHPPFSVPP VLSPLQRAQLLGGAQLQPCRM SPSQFARVDGFYGSPLAAMPYK SQAPMFRPDTTHLHPQHRRLLH QRQQQNRSQHRNLMGAGDRG RSSHQDHLKRDHYANLMGA RSSHQDHLKRDHYANLMG REKDWYSKIQMMQLQSTDPYL DDFYYQNYFEKLEKLSAAEEIQ GDGPKKERTKLITPQVAKLEHA YKPETIICQALCRALHINYLTKA FQOGPEYSASHITYK
14101	44469	А	14183	204	560	QGYWRGCTSSGGDGGCRGCW SPGKRR*GLSRTASGPPAAVAS PGRPRSPPSARTG*S*PGA*ARP CSRTW/PPPRCPLRPPSAARASA SRV/APL/GEPAPANTNIYCAAL THRATKSSTREF

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
14102	44470	Α	14184	59	1695	DMPWWPVLPWIKDSSPSSLEM
						NPDNTSGLRQKSVEAEPCSLW
		l			Ì	MVHRYTPQSTGEACTMAKAA
İ					1	GPSSP\GSCPPRPLGADPWPMSR
ł		ĺ			l	LSH*EC*KTGTTASGYRRHQET
		1				APPESSGSPVGLSAGLWLSWA
				l		WPVSPACLHPGPPLCP/GPPALG
				l		LGPRSPAAS/C/PPLPEAPPACSP
ł						FCP*GEISSSGPLEPVPOV*LPGD
1		l				LSEPPLGTSPLGRAPGLCSLLRA
		l				P/RPGAACPPASPSRPAMLLS*E
1		l				APRRPGPSGETSPSPSLAPSEASE
1		l				G\PPR\HHNGRDSEGAGTRTPGA
		l				SGSSKG/LGGQPAPHGPGGR*G
		ĺ				GGRYRGSAAGSVHSAPGGCSP
		1				HAPGCA/CDVRG/AKPSHFHSG
1						CQSG*GHQTSGGCPHLGLSPCS
1						TCLCAGQLQQQK/PGSSSGNL*
1		1				GQGHSSGDLGSAPPHLVPG\PG
					ł	PLTAGGHG/SLVGDSEWGSPAP
		l				GAVLGQNSSGPDPQRAGRWLC
		l				SRAEKSGCYTAPPLG*LSAQFQ
						EEKASWVPPGSGOLEDSHPDPY
		1				TALPLAAPPWGTGTGCPPAWP
						RL/EPDDCGPRHLSPPTCSTLPS
						WGGCAGS
14103	44471	A	14185	264	643	RCLRPGEGRSRQGRRSRTGRGK
		l				AEEGGARTRQQRRAALLPERS
1	:					ASQPGPRDSPRWRRRRQTTRGR
1		1				ARPASSGGT*GSSWLTLAWPPR
		l				EP*RLRSASAQG/CRRQRLQQR
		l				RRGLYQPHRPATCGPRAAD
14104	44472	A	14186	114	1039	QSQRLRPLADTPRRSNRPCSAA
		1				GLLFPQSFWSQELSQGAFLLPPS
						SPCLHCLPGTAAASRHQRSPRT
	ļ					RPSVQERRETHGLCAHRCEWR
		1				AGSPAHGP*AHRFVSRAGSPAH
1		ı				GP*AHRCISRAGTPARGPYSRR*
į .		l				VSEAGSPA/AVVSAPCSLSS*AC
ł		l				VTG\CSPAHSP*AHRCASRAASP
1		1				EHTSREDAEQAP*MQGLCPVH
						QEQPPSRKSVVTSPHIPPETDCR
1		1	1	l		LLPKNPCANKHQPTATGGRAV
1						PPEVQHVPPRSGMTLGSHQPLA
1				1		LQFQMQATDAARLGLSPQGQG
						SPPGPRSTLLLGCLSLFQLSLLR
		_				L

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	decenous, i-possible nucleoride misertion)
14105	44473	A	14187	1	1629	MIIYVENPKOSTSKCVRAYTFS
					İ	KVAGHKAIEKRLEKLAVDPHR
						DRAQMLKVENVQQAWQQWIN
						KLPPARREDEDVKEIRWMIEEL
						RVSYFAQQLDHTGVDRGQPGA
	1					STWSASLMCSVTOPKLKTRKP
					1	GFLLGLGERNCFSVYGAGRSW
l						GRGLGSCGFWLPPRTRIRAADV
		i		ļ		CLARWGWRAHRVRVLAGKNS
1				1		RAESMLWSFHKAAIEAARTNII
					1	QAQTRVEAAQATERRIAADIDD
1		1		l	ł	SELKAPRDGRVOYRVAKPGEV
1		1			ł	LAAGGRVLNMVDLSDVYMTFF
						LPTEQAGTLKLGGEARLILDAA
						PDLRIPATISFVASVAOFTPKTV
						ETSDERLKLMFRVKARIPPELL
İ		l				QQHLEYVKTGLPGVAWSFLYSI
1		l			ľ	NOTICLELDSIEAKLQALEATC
1		l				KSLEEKLDLVTNKOHSPIOVPM
		l				VAGSPLRTTOMCNKVR/CVNP*
		1				ATVVPPPVPQPTTQQYQGLDA
		1				GATTRVPRSLLRSEGSQTQKSP
					!	SCGSHSQDNSGGRKAVGARRD
1						RWWLGAGKGPDCRGAKCGLP
						LSGQKCPTLDPYQVVGCTSKTG
14106	44474	Α	14188	3	1010	GLOTOLVPLSSPVASLDNFSNL
						FGADLHVEKGQVPSSDQHLSQ
		1				RCHFPKAPPWSVCKALLPRSNP
		l				ATSLSGTSKPNSGGDFNSVTKS
		l				QPHCELNSFVLTGHSPVLWMM
		l				SEHDLADVVQIAVEDLSPDHP\
		l				VQSCGTVVVLENHVVTDEDEP
		l				ALKRORLEINCODPSIKSFLYSI
						NOTICLELDSIEAKLOALEATC
						KSLEEKLDLVTNKQHSPIQVPM
						VAGSPLGATQTCNKVRCVVPQ
		l				TTVILNNDRQN/DHCSQDGRPL
						EOOGTGFPGKMSLASO*HGTNP
		l				TFPCHCAESHENKSTSLSQGTV
	1	l			i	QHTLHLCGHRVEPGRHRPSSLR
						SGRAGWAMLNA
14107	44475	Α	14189	1	1710	
14108	44476	Α	14190	52	457	
14109	44477	C	14191	1	838	
14110	44478	A	14192	92	514	
14111	44479	В	14193	1	531	
14112	44480	Α	14194	1628	2008	
		_			· · · · · · · · · · · · · · · · · ·	

SEO ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14113	44481	A	14195	2	1616	CPPPRRPRPRAPPHHRPPAAGA
14113	44401	^	14123	ľ	1010	RTAAAGDRSRHGPASRVGTGP
						GOOMRKTYCRNSEPSDLDQGG
li						
						GSRGGPRRQSKCVRNWWVLGL TDFKNEAADPHDSGAQLASPSG
						FRTRAAGGAACQSRAMRSYSS
						ALGWSMGLGAVEOGVVLVGE
						AQAAQEPMEWVGGSGMVGCR
						SRAPPAGRQLRPDKKLSTAPVG
						RHRWRPSTPSAATGLGAKSLIA
						RGQQGRLAAPNKTDFKARAVK
						KDKEGHYIMIKGLVQKESITIINI
						YAPNTGAPKFTKQLPLDLRNKT
1				l		DSNTIIVGYFSTPLTALNRSSRQ
		l				KVNKETMDLNYTLEQMDLTDI
1 1						YRTFYPTTAEYIFYSSEHGTFSK
		1				TDRMIGHKTSLNKFKKIEIIPSTL
						PDHSGIKLEINSKKNLQNQTNT
						WKLNNLLLSDHWVNNEIKMEI
				1		LKFFKPNDNSDTTYQNLWDTA
1						KAVLTGKFIALNAYIKKSERAQ
					İ	VDNLRSHLKELEK*EQTKPKPS
		ŀ				RRKEIMKIRAELNEIKT/QKIQKI
						NKSQSWFCEKINKIGRPLARLS
		<u> </u>				KKGREKIQISSIRKK
14114	44482	A	14196	1097	1686	APRPLATREGPLPSNSGAQLASP
		l				SGSCTGLQVELPANSCTVRWQ
		l				VSALGWSMGPG/GPGAGGSAD
		l				QGGSGGTGAHGGVGRFRHGG
						AAGPEPCPMGEVAKAPARNRA
		l				QRQWAGTAGGPCASS/NSC*PG
		1				C*APYCPGWWGQPAAPSAGPP
						SPRPPRTRAGPQAPWAAPVPAH
						ASP\PHLPAS*GSPLRPGIQGSKT
		_				YQKSH
14115	44483	A	14197	647	1496	FLNLLSWQNLMLCGLKKKRKQ
1 1						FKDYEKKVKELNEERDKYRKG
						SVYIYTFGKEPTLNKVLDACDG
				l	i	KFQAIDFITPGTQYFMGCRWSC
						LPVLHCALAGLSPWVVDGTRR
			1	I		PGAVGSAGRGGSGGTGAHGGV
			1	I	l	GRFRHGGLQVRSPAHGEVLRPS
		1	1	I	l	EKSSAAQWAGTAGGPCASSTA
			1	1	l	ADLGAKPLTARGGGPAG\PPSA
				I	l	GPPSPRPPRTRAGP/PSTVGSPG/
		ĺ	1	I		SPPTPLPPHLPAS*GSP\SGLGQP
			1	I	l	RKGLPQCSGGLKGWLLKHRQS
		L	L			GCPGRGGAQSERGLRGLPARC

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14116	44484	Α	14198	1	560	MAALLLVGASW/LLLLVGAS WLLARVLAW/TYAFYDNCHRL QCFQQ/PPKRNCFGHLSLCGA/ TEEDMRLMEDL/PPLFRDVQL WLGSFYPVLSLVHFFTAPV LQASGG*PGFFTTMGLSFKRAK WQRLA/SEGSVRLEMFEHISLM TLDSLQKCIFSFDSHCQMVSPG SCTTSRGPRIPGALPTGGARAPE NGDPKEIE
14117	44485	A	14199	175	2225	IRWSSVSFASGMPSAPCCWGIPP QCPSAAQPTGPSWAGCRGLIP PNGQASPAS*VPPGAFPFLCIKQ PLWENLMTGVGKFSREAATLE GPTLETRIGVGPQCRQIADLPLE RTEGWRGDIGNPVRGVRQKRK UVAGETAASLLPCASGESCS VP*QARGPPVLFLPQ;WGRRWA KGCASRSCF*PDFRFSIIA*VPGA KGCASRSCF*PDFRFSIIA*VPGA KGCASRSCF*PDFRFSIIA*VPGA GCGGLQGGSSSAAQAL*PQLS GPRLPREK*GAPCWRGGPRCC PQNSAPETLEY*VEPRGTGAPGT THTGLPSPSQGCRGAEGWWGI GLARPRDPGOFSGVAPAGSAGFR ARTPPPASWAAKGRGRLAG GLARPRDPGOFSGVAPAGSAGFR RYPRASARAFGRGAGSAFPG RYPRASAFAGRGAGSAFPG RYPRASAFAGRGAGSAFPG RYPRASAFAGRGAGSAFPG GGAGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14118	44486	Α	14200	1	2850	MHPQERTAAATNAAIIITDGAA
						TOMGVTSTPAPGHPRDGYYOG
1						KDVSLLAHTHSFKISAAIEAKTT
						WVGMCRWGCKVALDVTFDVH
1						OGKPEGAGRNKQIPFIQDFPFAL
1						RSEKRPSLPATISGKTLPLNTKT
		l				CPQLLQCKREGHSLQCLIWLQ
						MVPQLGRLAVGGCHSNAWFPS
						KKAVIWEKFKHNLSGLVSDKG
l						GKRTGKGLGECLLMMRKMSP
		ŀ				ALEALTTFDSSLQHHFPRTFPVS
14110	44487	<u> </u>	14201	1	1570	ALEALNPGKPRQLVTPA MOSATSDAGSAPKKOKKVMTL
14119	44487	Α	14201	1	15/0	
1			i			QEKSGIAWYFFTAAQEWPNTPP
1						KLRTCTESMGCCLLDCCCTGQ
						GDWTRVTLYCRYPFILFGTNA
1						WVANFERPRMNANSLLASPTG
1						LSPYLRFGCLSCRLFYFKLTDL
		l				YKKTTLGSLPDAAMAFVNCHG
		l				AGGTVAVRTSRGHSRCHFGFG
						VSGTRYYHLTPQVVWERQQQK
						VDSLLGTQQMLEAQGREF/PGP
						CIPLPVPTNAPAAKTTPQPPSPD
						L*PPNYLFLCVDGHRPIC*EPKQ
						NKSDSFTSQA*SQDG*YCGCS/C
						RAPRPHGTSATIPMSSSCKRFQP
						SSPF/CKGPYHFRFLEERPLASRP
						SIHPSCGSYWLRLGCVAVGRTG
						/SRLNLOLHSASGLOALSKOR*R
						YCQSAGAGSPEPPRGGTR*CCR
					1	EAAPGCRAGTCPSPAANLPSLR
1						RK/PGTC*PQNPPGRCGSCGASA
						AVLAETPARIRRSFPRARQSCPR
						TWMAAGAPGSSRROGTKHVT*
						*SSTLEKOLDNPTATYFSVPTRK
						PVH*NQQKDL*VTG*LQIFHL
14120	44488	A	14202	11	356	VSGPPR*APWCPCLAAPTGSTS
14120	44400	l^	14202	l''	330	HGSTARDSSTACKVS/GGATCV/
			l			SGCGPGLGVLGRGLSSTRSCCM
1		1	l			
			l			PAG*CPSSKDTCEKSGGRTQPV
1			l			PPFEWAAQDASLVRRCRILAHR
		_			.=-	PCRGTS
14121	44489	С	14204	333	473	

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
_		<u></u>				
14122	44490	Α	14205	344	1035	QSAAQWFWWPGRSASLGGAK
	1			l .	\	GMQPPSLASWPHPRSIRCLRAP
				l		AP*QFTNATDPRFRSAPRRSGQ
	l	1		1		QQPRGSSGPAALASPSPRQPNR
ŀ	1	1		i .		AGALRTGGAGWATALDTWSC
		1		1		PRTRRRAPMGSCL*AAAQQCV
l	i .	1		l .		TGATWPPPSAIA*QSAAQWFW
	i .	1		i		WPGRSASLGGAKGMOPPSLAS
				1		WPHPRSIRCLRAPAPCSGPSASS
1		1				AAVQVACCCSLACCGPSRPASQ
	l	1		1		GHLDGTPITSPETCTTSLWSPOR
			ļ			
	1	1				RRAAGPPKVVDIPD*RGSRELPS
				l .		GRGAPNTTCIPYGGSPGAKWIE
				1		GCPAQ/RPSPPGLHPAMRASAL
				ı		LLMPFLQRRHRSATRLLLRPGS
l	1			l .		DCTGKK/GKGFLAGPETKRQKR
				1		LCPSPKTLGRALGQQEALPANV
l	1					RC
14123	44491	A	14206	41	292	
14124	44492	С	14207	480	1637	
14125	44493	Α	14208	3	410	SPPCGAPPPPVFCAPAPRAPAPP
		ĺ		ı		RAPPLAEAALPGFASAAGSAPA
	1			1		RSGPAGAG*GSRSRPRARASAS
				1		LAPASAGDHLGYPSATAAAAP
	1			i .		ASRPWLWRPSLNAARLGDPAC
	1					GWQASQWAAGSA*RAGSQWA
1				1		AGEARA
14126	44494	A	14209	1	1274	FFFFHPLRSLLGDLGLHSQERSA
14120	774274	l^	14207	ľ	12/4	PEIKPGALESTWSQARGAALRL
	1	1		l		AEAR*RAAPAAGTACSAGGAA
	1	1		1		SSPR*GSRATRRRRRGPTAPSRA
	1	1		i .		
	İ			1	l	ATAPGRGAAASAAPAARPART
l	1			1		ALMASARGPP\TAASPAQALTR
						SAPGHLPPMSELGAVTALCESC
	1	1				PVT/AAPRSGGGEAGGTLASEW
	1					GRTCRNYG/PAPGIWPRPPLSAA
		1				RPAAASPPARRIWSPPAPVHSA
						ASPIGLRVAAPCSSGSAVPLRSR
		1				RASPPCGAPPPPVFCAPAARAP
						APPRAPPLAEAALPGFASAAAC
			1	1		RGKLRLALQGGLGRSGSRSRQE
l	1	l	1	1		GPAAGS*LRTGKSWPSPPPGSA
		1	1	1		PLPLAQVLQALVEAGSRSRPRA
l	1	1		I		RASASLAPASAGDHLGYPSATA
l	1	l		1		AAAPASRPWLWRPSLNAARLG
l	1	l		1	l	DPACGWQASQWAAGSA*RAGS
		1			l	
1	11105	١.	14210	-	225	QWAAGEARG
14127	44495	A	14210	2	335	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop endnn, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14128	44496	Α	14211	344	1062	TRAEVTQQKLSRFASPPPVLRL
						RSYVSPPAAPTADPRWRPSWST
				l		SLRWVVKPNVSAYRIPPRASNR
				l		GYRASDWKLDYA*\WTGRLRIT
				l		SKGKTAYIKLEDKVSGELFAQA
						PVEQYPGIAVETVTDSSRYFVI\
				ł		RISDGTGRSAFIGIGFTDRGDAF
						DFNVSLQDHFKWVKQESEISKE
						SQEMDARPKLDLGFKEGQTIKL
						CIGLCSKPGTTAIQLGPVLNGIG
						RTLRTDLRNKNDLEGTNL
14129	44497	Α	14212	573	1017	TDQGVGSPHCPRFQQGPPRSPR
						FQQRSPHSPDSSRG/PPHSPRFQ
						QGSPHTPDSSSG/PPYSPDYSSG
		1				HPTP/HTSQQGIPC*DGHGWLPC
		1				GPVFLWSEAPHLNPGLGVWGQ
						PRAPALPGNSAESKLWPHDRPA
						RPETPGAGTAWAHQALRV
14130	44498	Α	14213	707	836	
14131	44499	Α	14214	724	927	SFMDFRFVLLFFHLVPNPK*RP
1.						VS/SPPGSKLLVPPWGPKT\PVPP
						HPSPPPPAHGEPGLPPPPPPPLPH
14132	44500	Α	14215	29	157	
14133	44501	Α	14216	I	363	
14134	44502	Α	14217	770	952	
14135	44503	Α	14218	1053	1995	KSSEPDLSHSFIFCSPNFPPSSSL
		l		i .		KRPH\FQSSILACVFCITADSSRA
						PCVLRVCDGFAGLTRSSAPSSK
		1				EKPCGFSLPEFTPLAVSFPQAAL
						CSLGTVSCLRPRPQLTPSWECP
1				ŀ		VLLAGAVCPGLPCAAIGCPWA
						GEEGCPGATSQLDQVSWPSLQE
						APEVLWPCLDSWGWFWAPSTV
		1			İ	IPLMASPRFGFMGRSQQPVLIN
			ļ	1		HLIKEPKAKFSNKNHSIRCCQG
						QRKWRRGNAAGMCCQPQIPGK
					i	GEVKTEFQPVSNRSVNYGCTM
						HFQETLWTPRLGLGDGDSNNN
						NNNNNNNRITANPGKEAEKPE
						HPHLLVKM
14136	44504	A	14219	482	833	THQGARGCTHARVSVREMRTR
						RGRWATGMRVPAPCSSVHTTA
1	1	1		l		VQRQQVADQLQTTAELSME/IL
	1					N/KWDQDPEAFVRKVATSYET
						WIS*YYPEDKAQSKQWLPRSGS
						GATLYSATNQYP
14137	44505	A	14220	195	367	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14138	44506	Α	14221	989	2210	LLVSCAILLSVSFLHSSSFVRSRS
						NRDSPACVDATYRGCSCQGEG
						VCGEACSCFPDLNQPQGVHAG
		l				EGVLPCSGWEDTFGKNTCLVS
ŀ		İ				YQQIPNKRRPCTCEECGKAFGQ
						RSHLVQHTSEKLYACQECGCTF
						SNNSSLVKHWHVHTGEKPYMC
						GHCGKCFRESSSLAKHQRVHT
1						GEKPYVC/GESTHLVQHW*FHT
1		l				GEKPFAC/HR/CSKAFADFSALL
						ACHGTYTGERPYECRVCCKAFS
1						PSLSLAEHIRCHTGEKLYACQE
İ						CGKAFSHSSSLSKHQQVHTECG
1			l			KAFSQGSYLTQHLKIHSGEKPYI
	ļ					RGEHPYACGKCGKTFSHSKFLT
	1		l			QHEQVRMGEKPFMCGDCGRAF
1	1		l			MQTSSLALHQRTHNGEKPYKW
	1	İ				NECGKSCIQMSHLTEYYQNLPE
						KGSKDPHTDATVHSPVMD
14139	44507	Α	14222	1	245	PPFRKQQAQHRNKTVLVSRSQP
l					1	VPVTVPE/PSA*RSTPEGPRRAG
ŀ						CDPRP*PCPAGPELPAQMKVLA
		ļ.,				IEQLLSLHFKKALLF
14140	44508	A	14223	477	947	LCSLSSLPPPPPPPPPPPPPPPPPPPPPPPPPPPPPPPP
						PFPPPSPVPLPPSPRSPPVSPPPPH
					i	SFQGRSPPSELGSSAEPWLRPGT
				1		WV*/PPPLTFSQQEAASSAQK*N
				1		CPGES\PQPVPVTVPERQPDAAP
						NHSLLLPGQQCFWCQGPHPPY
	L	_				WMGPVDASSSGFLSQSFY
14141	44509	В	14224	438	1437	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid scquence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop cudon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14142	44510	Α	14225	584	2866	WHFAKLSRNTQDHTRPTLGAA
					1	OAVPATGOPAAPGPAGAGPEW
						VPFLAVYKEQLVRNDSVLKVFF
					1	FPPIFYREGWVCCSVVESHLQE
						GFVWLFSLQL*PPKLRSLFSLSY
						CLSP\SDIEISRGQTPKAVDVLA
	ì					KEIGLLADEIEIYGKILVSSAGF
	ĺ					A*VVWEIL*NSLSPPELRITPTPL
					1	GEGKSTVTIGLVQAVSLGLCPS
	1			l		WQLYLVTPALQPGQQSETLFLK
	1					INK*KDILFL*FNLHLTGDIHVVI
						SCELSKFMDN*LLCGLIFVLCFF
						RGIILVTLCFLFLLKLLTFFLLNH
1					1	*KLGINKTDPIRTMPDTHQVAG
						PYLST\DD*SVLFQ*SPAPCGNG
1	ŀ				1	HPTGTVADHYLCLGW/YQAQF
	l	1				DIAVASEIMAVLALTDSLGREV
	ĺ	1				PRG*PLPG*YFVRNLMFFSHLPQ
					1	IDLGFHASVCLLGF*FAFSLL*G
						TPVFVHAGPFAADRDLCPYQFL
					1	LCQTHIYKIHVIMAEYFQRYKL
				İ		MMYAKQTSPLCDLEQAISLSGC
				1		VPCRHSHLKNEKFSCTGC*QYL
		1				PKLSLPNSDPLTLPFSADYGFKF
						ESNHANHSGDCQNLPLFLEDNS
1	1	1				TYSTCFPTVTEAGFGADIGMEK
		1				FFNIKCRDVVLSFRSVV*AFLLL
1		1				AV*LGRFGGRQYTGVWVLLFS
		1				FLQNIQLVADGV*MVSLSLKT*
	1	1				LNGYRCKTTNIFFTRTDTRAEID
	1					LVCELAKRAGAFDAVPCYHWS
		1				VGGKGSVDLARAVREAASKRS
14143	44511	Α	14226	3	587	YPASAGLMLQNFGVIGLRYHF
	1	1				AIHSPAAGGLLDGLHAVAAIQG
		ı				ITKIETTPNHQRAPAHWTLLTQ
		1				QAHLQPSPLHFNLPLTLCLMHC
	1	1			1	PTAIPHCFADARTWVNLPTSSLI
		1			l	GHKKENLKEFISGSLIVHEILEE
	1				1	VLQAEGDFQPFTRVTVHWGKG
	1				1	NDQTFRGLLDTGSELTLIPGDP
	1			1		KHHYGPPVKVG\AYGAQLL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14144	44512	Α	14227	2	1256	CHCGPP/VKVEAYGSQVLKGVL AQVQLTVGPVGPRTHPVVIFPV
						PECIIGIDMLSSRQNPHTGSLTG
						RVWTIMVRKAKWKPLELPLPR
				l		KIVNOKOYHIPEGIVEISATIKD
		ŀ		l		LKDAGVVIPTTSPFNSPIWPVOK
		i i				TDGSWRMTVGYCKLNOVVTPI
						AAAVPDVVSLLEQINTPPGTWY
				İ		AAIDLANDFFPIPVHKAHOKOF
						AFRWOGRQYTFTVLPQGRWEI
		1				NMTKIQGPSTSVKFLGVQWCG
			ŀ			ACQDIPSKVKDKLLHLVPPTTK
		1				K/EAQCLSGFRREHIPHL\PIYRV
		l	ļ			SRKAANFEWSPEQEKALQQVQ
		1				AAVQAAWPLGPYDPADPMVLE
		1		l		VSVADRDADWSCWQASI/GHK
		l				VGHAQQHSIIKWKWYIRDWAR
		l		l		ADPEGTTKGQGQRRWWQLAE
		1				RQDSRDREAAIGERQETAVGKT
						ARDGEAVCD
14145	44513	A	14228	155	531	GNLWSVDLRPGTPLRQNFRGTI
		1	ŀ	l		RQQHSRFTKNHCSQTPLLIPRQ
		l		l		TGSGVDLSKLQQTCS*GSCLVC
		1	ŀ			TIDLANAFFSIPVHKA/HQKQFA
		l	i			FSWQ\YTFTVLPRLTWLQPC*V
		_				PNLPAAETNTEPSNGT
14146	44514	A	14229	1	518	MTVDYCKLNQVVIPIAAAVSD
		1				VVSLLEQINTSPGTWYAAIDLA
1		l				NAFFSIPVHKAQQKQFAFSWQG
		1				QQYTFTVLPQWYINSPALCHNL
		1				IRRDLDCFSLPLDITLVHYIDDI
		l l		l		MLIGSTIKWVVHSS/DSIIKWKW
		1				YVHDWARAGPEGTTNGLAG*S GTCKKHEWKTGDKGIRGRG
14147	44515	A	14230	281	[140	VRVLSPVEKELKLWKNTHKLL
14147	44313	^	14230	201	1140	SYPTYGAAVTQLQNLTAMGVI
			ŀ	I		GSHGARGQVVALNRQRQGDL
		l				QPFTRVTVHWGKG/NMQIFGGL
		I		1		LDTGSELTLIPGDPKHHCGPPV
						KVGAYGGQVINGVLAQVQITV
l			ľ			GPOTHPVVISPVPECIIGIDILSS
		1	l			WQNPHIGSLTGIMVGKAKWKQ
1			l	1		LELPLPRKIVNQKPYCIPGGTVE
						ISATIKDLKDAGVVIFTTSLFNS
1			l		1	PIWPVQKTDGSWRMTVGYRRL
		1	l	1		NOVVTPIAAAVPDVVSLLEQIN
			l			TSPGTWYAAIDMANAFFSIPVH
14148	44516	В	14231	1	1521	
14149	44517	Ā	14232	3	202	
14150	44518	Ā	14233	2	367	

			1000			
SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	Sequence			sequence		[
14151	44519	A	14234	l .	553	MTVDYCKLNQVVIPIAAAVSE
						VVSLLEQINTSPGTWYAAIDL/
		1				NAFFSIPVHKAQQKQFAFSWQ
						QQYTFTVLPQWYINSPALCHN
		1				IRRDLDCFSLPLDITLVHYIDDI
				1		MLIG\PRQLLACY/WALVETEH
		1				TISHQVTMRPELPIMNWVLFD
		1		1		SSHKVGCAQQHSIIKWKWYVI
		_				DWARAGPEGT
14152	44520	A	14235	3	728	
14153	44521	A	14236	1	635	
14154	44522	A	14237	3	266	
14155	44523	A	14238	1 2	1245 837	CHOCKE AND COME IN COME
14156	44524	Α	14239	-	837	CHCGPP/VKVEAYGSQVLKGV
			1			AQVQLTVGPVGPRTHPVVIFP
						PECHGIDMLSSRQNPHTGSLTC
					1	RVWTIMVRKAKWKPLELPLPI
						KIVNQKQYHIPEGIVEISATIKI
				l .		LKDAGVVIPTTSPFNSPIWPVQ
					ł	TDGSWRMTVGYCKLNQVVTF
						AAAVPDVVSLLEQINTPPGTW
					}	AAIDLANDFFPIPVHKAHQKQ
						AFRWQGRQYTFTVLPQGRWE
		1			ł	NMTKIQGPSTSVKFLGVQWCG
		1				ACQDIPSKVKDKLLHLVPPTTI
14155	44525	١.	14240	1	533	K/EAQCLSGFRREHIPHL
14157	44526	A	14240	1	1043	MTVDYHKLNOVVTPIAAAVP
14130	44320	^	14241	'	1043	VVSFIEOINTSPGIWYAAIDLA
		l				VFFSIPVHKAHQKQSVFCWQC
		1				QEYTFTVLPQVHINCPALCHN
					i .	LRDFDHFSLPQDITLVHYIDDI
		1				LIG\PV*NMLLHLVPPTTKKET
						HLEVLFGFWROHIPHLATPSSE
						NGPMNKVTMMA/ANVGYAW
		1				QKHGLPLTKANLVMAIGECSV
		1	1			SPAAETNTEPSIWHHSSGQDSG
						VQESKGGNGSGTTHHHP**ST
		1				KIFASCSHDITFCWPRGLSSRG
				1		NAATRRHNNNFIKPEVKIAFW
		1	l			LWDPPTFKSTG*ERS*SVGWW
			l			D*HELPR*NQSTTPQGSVSIIPH
			1			GOPPHOLFGVAKARTFGIT
14159	44527	A	14242	1	336	OQI I IQLIO VAKAKI FOII
14160	44528	A	14243	3	1885	
14161	44529	B	14244	1	1423	
14101	44369	ъ.	17274	<u> '</u>	1423	L

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location uf first codon for peptide sequence	Nucleotide location of last cudum for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		L				
14162	44530	Α	14245	1	1361	MIISIDAEKAFDKIQQPFMLKTL
ļ						SKLEETASPSPVVATYTPQPML
		1		1		PSAFPPLSEEINPVLPETTVMAS
1		l			l	PEAVTRQDNVDSPQKPPPTPMF
				l		ASRPITRLKPRRAPSEEGIQRLK
						KIGMVEWISHFRPTPLSMEGPE HILLTNTLLNRYVKAAPASLKS
		l				
l				ŀ		PLTALLFMSDLTVGTTFSQLQN
						LNTMGIFGSSCDRSQVAALNHQ RQVPECKIVIDILNIWRIPHIGSL
						TGRVREIMVGKAKWKPLEOPP
		1			ŀ	PRKIVIQKQYHIPGAITEISATIE
		1		l	ļ	DLKETGVVILTTSPFKSSIWPLQ
						KTDGSWRMTVNYHKLNOVET
1		1				PIAAAVPDVVSLLEQINTSPGT
1		l				WYAAIDL VNVFFSIPV YKAHQK
		1				OFAFSWOGOOYTFTFLPQGYIN
1		1				SPDITLVHYIDDIMLIGSSEQEV
		l				ANTLOLLPARVASWGVPYDQL
		l				T/GGRED*GLLHRWSCRICRHH
		l				POGSWIQES
14163	44531	A	14246	1	1795	MRKCGKPQFKLGQTNKANSRI
						QEELIHSKSLIEQEGEKPVQFSA
		l			ĺ	FHRMWQPADSQCDIIDSADIWA
		l			İ	DPLVRHREIITGSGGINRRRTRG
		l			l .	GRGRGGEAAGGAENCGSREER
		1				ERAGVGTAVTQLQNLNTIGIIGS
		1				RGGRGQVAAINHQRQGGHSYC
						KGQQKQNSNQNSVTHVELWH
1						WLINHSVPRSEIDRKPTTFLLNL
1		1				YKQKTSRDLWPFTRVTLHRGK
		1				RNDQTFQGLLDTGSELMLIPED
		1				TKHHCGPPVKVEAYGGQVING
		1				VLAQIQLTVGPVGSGTHPVVIY
		1				PVPECIIGIGILSSWQNPHIGSLT
		ı				SRKTDGSWRMTVHYHKLNQM
		1				VTPIAAAIPDVVSLLEQVNTSPG
		1			!	SWYAAIDLANAFFFIPVHKAHQ
		l			1	KQFAFSWQGQQYTFTVLPQGE
1		i i				TLVNFSLPQDITLFHYIDDIMQI
		l		ŀ		GSSDQEVANTLDLLPRKSTTPS
		1	l	l		G/LYGFWRQHIS/HLGLLLTPIY*
		I	l		l	VTQKAA/SFEWGLEQEKALQQ
		ı				VQATVQASLPLGVYDPADPMV
		I	l	l		IEM/SLSDPSSHKVGCAQQHSII
		ı				KWKRYVCDQA*ASPEGTS*LY
		l	l			CTSFIMEKEEVC/LSLEQTLTLD
		I	l			MGLPILHAMLLWIHLWIHGLPY
			L		L	PSSWYSTQQCL

WO 01/075067

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14164	44532	Α	14247	55	1241	NRFVRAAPASLKRSAGIGLLCM
						PDLTVGIVVTQLKNLNAMGITG
						SELTLIPGDPK*HFGPPVKVGAF
						GGEIVNKYLAQLGLTDDPVGPR
						THPVAIFPVPECIIGIDILSSLQNP
						HIGSLTGRVRAIMKGKAIMKGY
					İ	YEGKGKAKWKILELPLPRKIIN
						QKQYRIPGGIVEMSATIEELKY
					1	AGVVIPIPTTFPLNSPIWPVRKT
						DGCWRMTVDYCELNQVVTPT
						AATVPDVVSLLEQINTSPGTWY
						ADIDLANAVFSIPVHKAHQKQF
						AFSWQGQQYTFTVLPQG\WTV
						NSPALGHNLIGRDLDRFSFPRDI
					1	TLLVHYVDDIMLIGSSEQEVAN
						ALDLLVFSDHLAIKWVMHSSIA
1	ł					SSSGSGICVIRLKKVLKAQPAP
İ						MASWGVLYDQLTEEEKIRA
14165	44533	В	14248	1	346	
	44534	A	14249	485	717	
14167	44535	Α	14250	1	345	
14168	44536	Α	14251	300	689	TKRAPCSPAAGSRGRARSLALL
					1	TLYHGHYPALPPGPTPAQQPGR
	ļ				ŀ	GLAEAAEPRGSEGGNGSNPCG
l	i					RA*DGRSRREGRGRLGGWRPC
i				[CEPQPWRQAHDPAGPDRVDGG
		1				ERRGAGVLRGQDGQDEKKKPK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14169	44537	A	14252		2430	MAVLQPSRHRLRPQLGLEPDN WHSQODEVSVATARAVYPSG RGAAWMGPSGNIGGSGLVSHE CGEGTGAAGSTQLQEGMHFCR SQRWAFLTALCEVYTGSLQDT VQWLQLCQWSAQGQDDHTLIE WLMGATWTVWNDAREIPETG TQDAWLCGSWOVDDQRPHV ELAHCSPTNVQRVLALVDTGA DCSLVFGRPYRKPDGTWQMTV DDWELNKVTPPLHAAVPSIMD LTDHLTMELGQYHFVDYVHSP TURGLVAMELAAWKCPKGVL LFHYIDDIMLTSDSLADLKVAV WPGLSAKFLGVIWLSKTKAIPE AIIEKIQAYPRPTTMRQLQTFVG WPGLSAKFLGVIWLSKTKAIPE AIIEKIQAYPRPTTMRQLQTFVG LEYWRPFVPHLAQMIKPLYGL SKKGATWDWDNEAGTDLLAA KWALQQTQALQVIDQGCPFELD VHYTTDGFGWVLWQKMCFGR TPVGFWSQLWKGAELQYSLIQ VHYTTDGFGWVLWQKMCFGT TPVGFWSQLWKGAELQYSLIQ AKRGAYLEQQSMLTTTPLAAE LQEALGPTVLMEDKAMGPEAPL DPEPSLKKGYPLYDGAWTTTPLAAE LQEALGPTVLMEDKAMGPEAPL DPEPSSQLAKAWTPAVIGOPTTT TI*FDTGCGQSTQWAELRVVE MDILEACQXCPACAQYLRQR MDILEACQXCPACAQYLRQR MDILEACQXCPACAQYLRQR MDILEACQXCPACAQYLRQR MDILEACQXCPACAQYLRQR MDILEACQXCPACAQAYLRQR MDILEACQXCPACAQAYLRQR MDILEACQXCPACAQAYLRQR
14170	44538	A	14253	2	239	DYAEPLPKLQGYTYALMAADI
14171	44539	Α	14254	3	211	VVYYSPNEVKVVAEGFDSANG INISPDDKYGPL*NIFIFLQLWC WPNSLLCFLCQCENSFLDIYGIL
14172	44540	В	14255	13	87	
14173	44541	A	14256	42	854	VLVFLTAALFILPTFSNSMMILQ VSGGPWT/VLIERGVLLGVAPP PSLPALVPENSVYQERQECYAFN GTQRVVDGLIYNREEVYHFDS AVGEFLAWMELGRPIGEYFNSQ KDFMERKRADQLPDCRDPLMP D°4VWISPLGQQVPNLTAQKTL QHLPSKKEPQQHPDLLVYHVT DFYPDSIQVRCFLNGQEETAGV VSTNLIRNGDWTFQILEMLENT PQQDRT*AGVQLLEVTLNLSLF FLEAQSDSVQSKMLTGARGLFI NYLQLNLCTP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \mpossible nucleotide insertion)
14174	44542	A	14257	3	571	QGRATPENYLFQGRQECYAFN
						GTORFLERYIYNREEFARFDSD
	1			i		VGEFRAVTELGRPAAEYWNSO
	1	ı				KDILEEKRAVPDRMCRHNYEL
		1				GGPMTLORRVOPRVNVSPSKK
						GPLOHHNLLVCHVTDFYPGSIO
						VRWFLNGOEETAG\VVSTNLIL
	1					NGDWTFQIVVMLEMTPQ/QGE
						DVYTCQVEHTSLDSPVTVEW
14175	44543	A	14258	1	1677	
14176	44544	Α	14259	2	430	
14177	44545	A	14260	821	1428	QVTDVQDFPLKCKIQWRRTVR
						GDLQ/PFTRVTVHWGKGNDQT
						FOGLLDTGSELTLIPGDPKRHC
						GPPVKIGAYGGQIINGVLAQVQ
		1				LTVDAVGPWTHPVV/I/SPVPECI
						IGIDILSSWO/NSHIGSLTGRVRA
						TMVGKAKWKPLELPLPRKIVN
		l				OKOYHIPGGIAEIGATIKDLKDT
						G\VIPTSPFNSPI*PMQKTDGSW
	1					RMTVDYL
14178	44546	Α	14261	507	910	
14179	44547	A	14262	1	2898	
14180	44548	Α	14263	1	2580	
14181	44549	В	14264	159	2657	
14182	44550	A	14265	142	377	NHLLRRQQWQTPFPPPSWS/SPR
	l					SSSDCCGSSENFKPVDLSLLGSV
	ŀ					GVGPTEPDHLDPWLQPPFQGSE
						RFCLAGVPGATG
14183	44551	Α	14266	363	638	EVWKESSGWHLAGAPLGQSFQ
	l					RKEQAVIFAVLQPLLVIPKQTGS
						GVDLQQTPAHLQQRGLSGCLQ
						SLCSLTSS*PQRHPQNSSLVSNC
						vvs
14184	44552	В	14267	1	2424	
14185	44553	В	14268	446	623	
14186	44554	A	14269	245	690	APSRSTWGSSSTAARRNLKTTR
		1	l			KLRTPRTPPHQKPRNPHHGPPR
						TPSG*TPRNPDHRPPPG\PQVGQ
					1	PGCWAPSQGPRGWGQLGC*RE
		1	1		I	EVPGGGHAGRHSTPVGG/DFFL
		1				QAGPWPPDASLPTNPFVLRGAS
	l					TPPHPLSENNKDKFLLQK
14187	44555	Α	14270	223	365	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14188	44556	A	14271	<u> </u>	687	MSEYIRVTEDENDEPIEIPSEDD
				[GTVLLSTVTAQFPGACGLRCRN
		1				PVSQCMRGVRLVEGILHAPDA
		l				GWRNPVYVVNYPKDNKRKMV
		1				LGLPWKTTEQDLKEYCSTFGD
		1				VLMVRQSQDEPLRSRKVFVGR
		l				RTEDMTEDELWEFFSQYGDVM
		1				DVFVPKPFRAFAFVPFADDQIA
		l				RSLCGEDSMKGISVHISNAEPK
		1	ŀ			RNSNRQ/LRSGRFGGNPGGFGN
						QGGFESYQRCERKAV
14189	44557	ا ـ	14272	1	833	QGGFESTQRCERRAV
14190	44558	A A	14273	3	447	RGTAGGOPGNGPALAOSP*WC
14190	44338	Ι^	142/3	ľ	447	RLGSGGT*GKSPGTSQGRALPW
	ľ	1	ŀ			CSC/SVSTWATAAPGSQQRASP
		1	ŀ			
		1	ŀ			CSWAQPGPLASLRLLMPPRLR*
		l				GHLGHLAWLQLFAAPCPVIHV
		1	İ			AVMWSLLGIALRKVHGDWAPL
		Ļ.			1000	AILSLVVCASNKLNLKKKKK
14191	44559	Α	14274	988	1803	RAARVPRAPLEAVAHRWASAS
		l				EDDGTVLLSTVTAQFPGACGLR
		l				CRNPVSQCMRGVRLVEGILHAP
1		1				DAGWRNPVYVVNFGDVLMVR
	1	ł				QSQDEPLRSRKVFVGRRTEDM
ł		1				TEDELWEFFSQYGDVMDVFVP
		ļ				KPFRAFAFVPFADDQIARSLCG
		l				EDSMKGISVHISNAEPKRNSNR
		1		1		Q/LRSGRFGGNPGGRLQGTRYS
	1	l			i	PAPAPGSPGRAQGLG\AG*SRPS
						GSWLSGHLQRPEQSVGRAPAA
	1					PRHAVRRSLRRLAALCRRTRAR
		L.				CRRAQLHVSPQRRRH
14192	44560	A	14275	3	184	
14193	44561	Α	14276	1	352	KAAMVRFKHRYLLCELVSDDP
			İ		1	RCRLSLDDRVLSSLVRDTIARV
1		1				HGTFGAAAC\SIGFAGTIRTCQK
ł						FLIQYNRRQLLILLQNCTDEGER
1						EAIQKSVTRSCLL*EEEESCEEA
						AKAME
14194	44562	Α	14277	1	456	PTRAVEAAMVRFMHRYLLCEQ
						VCDDLRCRLSLDDRVLSSLVRD
		1				TIARVHGTFCAIACSIALALARA
					1	LNEEGNPEGWRDVAGASAGGR
						KKGHECPSPCTVFCTL/RYPNA
						YTVIGLLRCKKELYQVVWSVL
						YLMTYLENKGHRYPCFFNTLH
14195	44563	В	14278	34	267	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14196	44564	A	14279	3	395	SAEVGAAETTLTELRRTVQSLEI DLDSMRNLKASLENSL\GILLHL
						ESELAQTRAEGQRQAQEYEAL LNIKVKLEAEIATYRRLLEDGE DFNLGDALDSSNSMQTIQKTTT RRIVDGKVVSETNDTKVLRH
14197	44565	A	14280	2	1044	SSQMPSPQEGMCGKACTPALS QADSLCPLLRLASEVEGYWQM ACAEPYVGSENNDHLSYWSLQ HGSTILASMSFTTCSAFTNYWS PGSVQVPSYGTGYSHAASVY AGLGGGSGSRISVSHATSFGGLA GMGGIQNEKETMQSLRDRLAS YLDRVAGLETENWKLESKIQEH EKRNET-ETEIEALKKELLFM KKNNEEEAKGLQAOJASSGLTV EVDAPKS/QDLAKNMADSRAQ EVDAPKS/QDLAKNMADSRAQ ESTRVYTTQSAKVGAAEMTLT ELRHRVQSLEIDLDSTRNLKAS LENSLREVEARYALQMEPSLLF
14198	44566	В	14281	247	857	SKIIFGRVWEISYLHVFEGKQV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14199	44567	A	14282	1	2104	LRGSELAGGVAAPDPSSRARAP GROGGSEPPRRS'GRGRAPPRA HTPLDDFPPGRAALSPGGLSS RRPDPGSAGRPLLCLRPPLHSP PPPRCCQAPAFOSLDSTHLLALR SSGEEKL WGPRRAPAPP SOR RKPDPARFGLAAAVPLRPV*AS D*GAAQVPPSPLRDGGQPGGR EGYGRRTPHRAAAWLLGE*AA DAAAFHWDGGRPPAAPARLLP QSPGDPTPAGEQIASRH*LCEN PETQKLRH*TSERRDGHREEH TGTAGVPRSRPATQRHAVPAG GLQPHRMLPALSSGAASGGEA EHGQLSGRGEEDCPWPQD AGLQGHFRGESPRWPPCLGDG EN*PGPVQAEFSGCDAVSESS EN*PGPVQAEFSGCDPAVSESS PATCG/RGGEFGVSSPKTLQPA ARDATRPQLLPRGRLPALSAEK HPDASSPWREPQAPRFSRCLH UPQ*MK*YEMTSPRAPTPSG RGCGPGPLSPRTPAAGRAGRGPR RGCWPRPVATHFGSPGQPPAL LPQ*MK*YEMTSPARAPTPSG HIGALSSAGVC*LQTTFE*IN **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* **THTWYHSEPTD*MGAEH*Y* ***THTWYHSEPTD*MGAEH*Y* ***THTWYHSEPTD*MGAEH*Y* ***THTWYHSEPTD*MGAEH*Y* ****THTWYHSEPTD*MGAEH*Y* *****THTWYHSEPTD*MGAEH*Y* ******THTWYHSEPTD*MGAEH*Y* **********************************
						GGEVIS*QQKGGGRAEHGRPTV QGP/SHGNGPHAV*PHAQPFWH PWGSILEVPYLTRPSGHHRIEH
14200	44568	В	14283	41	4224	
14201	44569	В	14284	1	1332	
14202	44570	В	14285	140	215	
14203	44571	A	14286	234	657	VQQPGRGLDLSTDGPGGRSQV GLIWSCCCLH*AASGEPGGRCP GSGAPGPAGSALEFRARDGVP! GVGGPSWESHSPAAATPPPAEC RGPGPTPSPAPGEAAPEDREDG AAAPGRAEPASIVAPADGSQGQ VLATQAGALGA

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14204	44572	IA	14287	134	789	PDSSQAPSRNSTRPAASTRKSSS
14204	44372	Ι^	14287	134	/69	PPAARS\AGNELGSSRGDADPPP
		ı		i		
		1				QNPTGPAPQSAEVPRCD/SSSA
		1		i	i	WASNGSSISPSFFSLRPWPSST/S KVPGAGHHTPPEGAGGASAR\S
		1		ļ.		SSLAPSAGAPVPS\A*RSQLGTW
		1				
		ı		ł		LSVLRPRSL*QSRVRSPSSHVSS PGSMQAPNSPSSFQPGMNGAWI
		1		i		WWL*THIPLARWPGPPGILVITC
		1		l		PTVTWIKWSFICSFHPT
14205	44573	A	14288	465	1541	GPPKRKLISNFSKVSGYKINVO
14203	44373	^	14200	403	1341	KS*AFLYSKK/QTESQIMSELPF
		1				TIASKRIKYLGIQLTRDMKDLF
		l				KENYKPLLNEIKEDTNIWKNIP
						CSWIGRINIVKIAILPKVIYRFNA
		l				
		l				IPIKLPMTSFTELEKTTLKFIWN
1		l				QKRAHIAKSILSQKNKAGGITLP
		l				DLNLYYKATVTKTAWYWYQN
	1					RDIDQWNRTEPSEIIPHIYNHLIF
		l				DKPDKNKQWGNDSLFNKWCW
		l				ENWLSICRRLKLDPFLTPYKKIN
						SRWIKDLNVRPKTIKTLEENLG
		l				NTIEDIGMGKDFMTKTPKAMK
		l				TKAKIEKWDLIKLKSFCTAKET
		l				TIRSFGSEQKSHTLVRATRLVSD
1				l		HSQSFEAWLQFLGDTRFGTQKL RDSD
14206	44574		14289	1921	3199	GYLAAGQVTSRIHTGRRHFSGP
14206	44574	A	14289	1921	3199	ARCPCVSCCRRPGC*PQPAVRP
		1		ŀ		SG*HCRKTAPSHGPSIWOTR**P
		1				GSWPNLRPG*GWRWLRAOPRG
1		l				RGIPVGCRSNLQLHPTKNPGSH
		l				*GSGSPWQCPGVLL*ASGRQPG
		1				
İ		l		l		SPSGGCT*SSGKAGLEPQWHPA
		l				IPGHCRNKPAAPASWERAPRPR
		İ				AHALACGWWRYSSC*SWRS*S
						SPETPGPAQV*CHRPCPAAWW
		1				RCHS**TAQFL*D*FS*FLQTIGE
l		1				PCICPKPISVPH*PMKQSH\RGIG
		1		l		GCLQYSSHPGIESSLGMSRGF/D
		1				RRDL*TLQIGS*RHPRAGPEGA
		1				AL*CLAGESSLRTPSAPCPGSPH
1		1	1	l		TPGSSASGNTSPV**ETGCGSRQ
		ı				GSARLVLESLLSR**CLSGPSRP
		I		İ		SCLLRGISWQCGTAGRLSSSGPP
		l	1	l		AASGAGADAAPGGFPRLPCLEF
		<u>_</u>			ra.	GTV
14207	44575	A	14290	3	626	L

SEQ ID	ISFO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1.00	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	,	1		sequence		
ŀ				· .		
14208	44576	Α	14291	1	1065	MRGSCERSGEDEEQKEEAMVA
14200	1.570	ľ`		ľ	1.000	CGRLSGVPEAEOGPEANWDSD
İ	ľ		ļ.		 	LETEGTDGLGELVRDTLYLRSC
		ŀ				
		ŀ	l		l	RAHSVVPISCFLRQGSAQELNL
			l			RHRGLGPQGARALASSLSSNPY
		ŀ	1			VKRLDLRDNGLCGAGAEALAG
						AL/QQKQQHP*CGPVGEPAGSG
			l			RSPGP/RVPPSQ/CNQAMRKMQ
		l				LSGNGLEEQAAQHLAELLLAH
	l	l			ŀ	TDLKSLDLSYNQLNDQAGPAPS
	l	ŀ	l		l	LRAFPOANIFLKVLDISYNGFG
			1			DPGASAVGEALKANNVLEELN
	1	1	1		ł	MSNNRISAMGALSLGLGLRVN
		ł	l			li .
1		l				QTLRILVLCVRVSQVSRNPMRS
			l		1	EGCFGLLKSVQDNPASALELLD
						FSDIQVNAEFDGLASSVRGILPG
						LGIKDWRLQSGV
14209	44577	Α	14292	1	416	
14210	44578	Α	14293	2	384	ELWVVTSFMAYGSAKDLICTH
		l				FMDGMNELAIAYILQGVLKAL
	1	l				DYIHHMGYVHR/DLQGYDAKS
	1	l				DIYSVGITACELANGHVPFKDM
	1	l				PATOMLLEKLNGTVPCLLDTST
		l				IPAEELTMSPSRSVANSGLMTA
		_	. 1001	1.42	151	IFAEEL INISTSKS VANSOLIVITA
14211	44579	С	14294	142	474	
14212	44580	Α	14295	1	1470	
14213	44581	С	14296	91	471	
14214	44582	С	14297	195	480	
14215	44583	Α	14298	121	1130	SRSTQYELMSFLTNDASSESIAS
		1				FSKQEVMSSFLPEGGWYELLTV
						IGKGFEDLMTVNLARYKPTGE
		1				YVTVRRINLEACSNEMVTFLOG
		1				ELHVYKLFNHPNIVPYRATFIA
		ł				DNELWVVTSFMAYGSAKDLIC
		1				THEMDGMNELAIAYILOGVLK
	l				İ	
		l				ALDYIHHMGYVHRSVKASHILI
		1			l	SVDGKVYLSGLRSNLSMISHGQ
1		1	l	I	I	RQRVVHDFPKYSVKVLPWLSP
1						EVLQQNLQGYDAKSDIYSVGIT
1		1	l	I	l	ACELANGHVPFKDMPATQP/M
1		1	l	I	I	VLVLSSPDAARETERHSALPVG
1		l	l	I		YOHHPR*GADHEPFALSGOLW
1		1				PE*OPDHOHPPALQR*LALPPLP
1		1	l	I		PNLLPPLPPLC
14016	44504	-	14200	704	412	I MELLITERE
14216	44584	A	14299	304	412	-
14217	44585	A	14300	2	350	
14218	44586	Α	14301	191	299	HRPATAVLHVPTRSPPSPGSRPS
1	I		l	1		*PPPAAALLWTPA

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14219	44587	A	14302	537	5523	LARYYFFMHODRQGVARNILL CQKEKRR*YTTKCEDNTCHWT GRTSS*PRSVISNLNSQCGPRLW PPVTIIRQANIKTGEKAH*TSKL LPPVQLKFPARDDTLPWPGYIP* QLRRISPR*SILSMLPRKVVPCL *QKP*TVCWKF*EPLSYIHIBLY LKMMIPMPMTLLGA*CLTQAL K*SRMERKGRAAQRSSTLGC QSPRGSPGGFGPPGPGAGKQ QSNSVQTSRLLARTLI*SFSSST VKDKELARNTAP
14220	44588	A	14303	2868	3095	PGWAGGLFGPSAGMRAWATK PSLTPPAPAGQRCPQGHRG*SG PGIYAVQPPSVPAWAAHATTQ ARLCHPRSPPLR
14221	44589	A	14304	693	821	SDRQT*RQEKRLTRRQSCCLPC N*SFQPETTLFHGPDTFHDN
14222	44590	Α	14305	338	645	_
14223	44591	В	14306	181	480	
14224	44592	Α	14307	155	543	
14225	44593	В	14308	1	1260	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14226	44594	A	14309	Ti	2025	VINIINAAQESSPMPVTEALDRV
		1				LEILRTTELYSPQFGAKDDDPH
1		1			İ	ANDLVGGLMSDTGLEIIKDGEE
						EGESRTAQQYPGLSESTRQPWR
						RSRTPRTTWGWKATEQQCPDF
1						PPPGPHPHLELLIFHCEGQGTGQ
						KHSTSYPCTLMSEKRGLMNVFF
1						AKYTGPAMRKSSGLTPSLLHLY
		ŀ				CADPLHSKPLFNCPPPTTSKTEI
1		1				LGSCNCSPSESVRGAEKGDOEL
		1				LANIKOAEKHEKNHPEVTVAM
		1				ALTDIDLQLQFSMSQPEALLLL
l		ľ		i		AAGPADHLLLQLYSGHLQSAT
1		ŀ				GIYGEGRASSATLEDLESQYQE
		1				LAVALDSSSTIISQLTENINSLVH
		1			ł	TLKEEKHEIHQVQKLGRNLFKL
i		1				KNOTAEPLAPEPPARPSKVEOL
		1				QDETNHLRKELESVGRQLQAE
1		1				VENNOMLSLLNRRQEERLOEQ
1						EGWOEEOERLLDKNNENKSAL
1						QLEQQVKELKL\ENLEATSQ*K
	1	ŀ				QQL/TAQLSLMALPGEGDGGG
					ŀ	HLDSEQEAPRPMPNIPGDLESR
						EAMVAFFNSAGANAQEEQRVC
	ì					COPLAHPVASSOKKPEVAAPAP
						ETGGESVCGETHOALOGAMEK
						LQESTSARGQCQRRSTGRGGHR
		l				QAGPGPGGDEEMLCTCTQVSS
						AGRGMGMLTVVLVVSLMMLS
		1				ASFSTWCMRSISSCHCAYSAHS
		l				SISVFWFPREYILASGSEMMNS
14227	44595	A	14310	3	564	SLRVSLLKAVVDRAGSRILSYIA
		1				GLGVDLGFGGGSRVAGGIADL
1		1				GRSGGGKATRSRRGSRAGGGR
1	İ					LRSRHHEEALQLRETPGPGAPE\
						PMDOEYAGPGYDIRDWELRKI
		1				HRAAIKGDAAG/VWSAAMTRR
		ı				FRDLDARDRKDRTVLHLTCAH
		l				GRVEVVTLLLSRRCQI\DICD\N*
		l				NRTPLMKAVHCQEEA
14228	44596	A	14311	515	778	GQAQGTGDREGAVLFTARAPD
1.1220	1.1020	ļ'`			1	QCRP*SCSSSRPTASSCPLGHSC
		l				SPGPLCTHLPLQGPPQAPTVPG
	1		1			ATATCWFSWNQCRCRAVSSYP
14229	44597	A	14312	411	608	EAEIP*TGLPGHSRPPGPLCTHL
1.722	1.757	ľ.	1.75.2	1		PLOGPLGAAAWVPEAGLGQVP
1		1			1	TEAAGAARAGCSAHKPWKRRK
14230	44598	A	14313	2	338	
14231	44599	A	14314	1	390	
. 7201	1.1000	<u> </u>	1	·	J	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14232	44600	A	14315	237	488	PWKTLDSSEAEILRGVSESSAV TLHCMEGQAVGPRQVKRIVHC *ASSA*NPHAMLLLQNR*LALC TSVPQGAAVLCGGWDPGN
14233	44601	A	14316	358	616	
14234	44602	A	14317	13	1155	IKLPPGCRRRGGHIRHIRVRG RAFPLAPRRRLPAPEGPRARP PASRFLKNTMSNGYEDHMAED CRGDIGRTNLIVNYLPQNMTQD EIRSLFSSIGEVESALLIROKVP GHSLGYGFVNYVTAKDAERAI NTLNGLILQSKTIKVSYARPSS EVIKDANLYISGLPRTMTQKDV EDMFISRFGRIINSRVLVOQTTG LSRGVAFIRPDKRSEAERAITSF NGHKPPGSSEPITVKFAANPNQ NKNVALLSQLYHSPARRFGGF VHIQAQRRFSPMGVDHMSGL SGYNVPGNASSGWCIFIYNLGQ DADEGIL WQMFGPFGAVTNVK VIRDPNTNKCKGFGFVTMTNY EEAAMAIASILNGYRLGDKILQV SFKTKKSHK

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ŀ		ŀ		sequence		
		<u> </u>		<u> </u>		
14235	44603	A	14318	1	2340	MFVWRNVEGHSAAVFSWYSIP
						FLTPPCSHTRPSNLPVTQWPPTR
l		1				ENNLPSWQPLLMSVHQAQSLS
		l				ALRKEQDSSSEKDGRSPNKSDK
1		l			1	DHIRWPMSGAHDLQQAAPGPG
1						GAHQGHPNQDNRTVSQMLSER
1		l				WYTLGPNEMQKYNLAFQNVE
		1			1	ARIRYVIVRGNEQGFFRMHHLR
		l				GVSSLQLGRRRPGPGTYRLEVV
	l					DMITIDARYRCSGLRSPPDPDH
	l	1				LKSVDLHRNEHFFLQKARWEIH
	i .	l				LVLQATPSQIQLLHVAFVLAKQ
						PLRCRKQTLFLISVVGTKEKNV
1		1				ASEGRNPVRPLRGHRWLPASPP
1		1				AAEHFAGGADRKSRLSRAPRSA
1					ŀ	ATATPANEQRAVADVGQWAR
					ì	ARGGRVRAAGRSGSCALRRSR
		1				CRRRRRHRRHRYRGRAEPLA
					ŀ	PRRAASGOPGKLRARRGINCNA
		1				
						SSIPMLQVVRIKSISKHCQMAR
					l .	VAGEGSLPLSHAQLTHCVFGN
		l				KQVQRTIAYLPPQQQGVQGFL
						KNTMSNGYEDHMAEDCRGDIG
					1	RTNLIVNYLPQNMTQDELRSLF
					ì	SSIGEVESAKLIRDKVAVGLLSL
					1	CVPIRNVEIVASTSWWSLPLAP
1					ŀ	YVPRGIRRSQPSTLMDDLSCRS
1		1				HQAAAPFTASVLLYISGRVKAH
		l				TAAQGFNHIPEEGVSQEAGSG
		1				WSQAWKDGWLTNVFMFSCTG
						HSLGYGFVNYVTAKDAERAIN
						TLNGLR/PPVKNH*GRPLQLGFL
14236	44604	Α	14319	701	1050	LIVHFCLQVGCGVGNTVFPILQ
					i	TN/NTNSEYDPSRCFAFVHDLC
					İ	DEEKSYPVPKSSLDIIILIFVLSAI
						VPDKMQKAINRLSRLLKPGGM
						MLLRDYGRYDMAQLRFKKGIL
						KVLNPNH
14237	44605	A	14320	701	1519	LIVHFCLQVGCGVGNTVFPILQ
1		ľ		1	'	TNNDPGLFVYCCDFSSTAIELV
1	1	1	1			OTNSEYDPSRCFAFVHDLCDEE
1	1	l	1			KSYPVPKGSLDHILIFVLSAIVP
1		l	1			DKMQKAINRLSRLLKPGGMVL
	1	1				LRDYGRYDMAQLRFKKGQCLS
	1	1		1		, ,
						GNFYVRGDGTRVYLCHTRCNW
1		1				TRFSPTAGLEKPQKP\VDRRLQ
1	1	1			1	VNRGKTTDNVPGLDSVQILQAP
	i					SVQHQLRGYLLPTRCKPVVVS
1	l	1				GLFLKKKIVALGVV/RCL*SQPL
		1				RRLRRGGSIE/HQQSNLGKIVRD
		<u></u>		L		PVSESNDKIKRI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=pussible nucleotide deletion, \=possible nucleotide insertion)
14238	44606	A	14321	756	1334	PGKFVLMSFLDPOPPTRILEVCG GVGNTVPPIFTNRTMTQDSLVY CCDFSSTABELVQTINSEVDPSS VFAAFVQGPV**RESVTQCPRAV LDHILIFVLSAIVPDKMQKAINR LSRLLKPGGMMLRDYGRYD MAQLRFKKEELDTLFTTAGLEK VQNLVDRRIQVNRGKQLTMY RVWIQCKYCKPLLSSTS
14239	44607	A	14322	249	474	VYLLIVLAVLYTNNRQTESQIM SELPFTIASKRIKYLGIQL\TRDV KDLFKDNYIPLLKEI*EDTSKW KSIPCSWI
14240	44608	Α	14323	1	5796	
14241	44609	Α	14324	1	2721	
14242	44610	Α	14325	1	3438	
14243	44611	Α	14326	1	3306	
14244	44612	Α	14327	1	2364	
14245	44613	Α	14328	1	2091	
14246	44614	Α	14329	1	2334	
14247	44615	Α	14330	1	1986	
14248	44616	В	14331	1	2559	
14249	44617	Α	14332	1	1671	
14250	44618	Α	14333	1	1845	
14251	44619	В	14334	1	2055	
14252	44620	В	14335	1	1554	
14253	44621	Α	14336	1	2559	
14254	44622	Α	14337	1	1714	
14255	44623	Α	14338	1537	2204	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14256	44624	A	14339	2004	3751	YQSLAETQQQQKKNFRPISLMN
		1				IDAKILNKILANRIQQHIQKLIH
	İ	1				HDQEGFIPGMQGWFNIHKSINV
		1		l		IQHINRTKDKNHMIISVDAEKA
		1				FDKVQQHFMLKTLNKLDIDGL
		1			l	EVLARAIRQEKEIKGIQSGKEEV
		1				KLSLFADDMIAYLENPIVSAQN
		i				LLKLISNFNKVSGYKINVQKS\Q
		1				AFLYTNNRQTESQIMSELP\FTI
		1				ASKRIKYLGI\KLTRGVKDLFRE
		1				NYKL\PLIKEIREDTNKWKNIPC
	l					SWIGRMNIMKMAIL\PKVN\YRF
		1				NAIPIKLPMTFFIGLGKILLLSSY
				i		GTKKRAHIAKTILRQKNKAGGI
ł		l				TLPDFKLYYKATVAKTAWYW
						YQNRDIDQWNRTEPSEIMPHIY
		l		ļ		NYLIFDKPEKNKQWGKDSLFN
		l	1			KWCWENWLAICRKLKLDPFLT
						PYRKINSRWIKDLNVRPKTIKTL
		1			1	EENLGITIQDIGVGKDFMSKTPK
		1				AMATKAKIDKWDLIKLKSFCT
						AKETTIRVNRQPTKWENIFATY
		l		l	1	SSDKGLISRIYNELKQIYKKKTN
1		1		1		NPIKKWVKDMNRHFSKEDIYA
1	1			l		AKKYMKKCSSSLAIREMQIKTT
						MRYHLTPVRMAIIKKSGNNRC
						WRGCGEIGTL
14257	44625	Α	14340	519	1286	LNLEERAFRHKSSEHLPHLKVH
		1			1	FRHSSCGIKTKSLLHVSPPTTLIF
				l		PYPPLWGNPSN*ARPPCSVHSG
1	1	l				PYPQGLASIIAHGSPGNVRGVH
		l				V*INDSEGGISRPSTASHKETPR
		l				PPTPARHADLAATRPRSLTTSPS
		1	ĺ	1		\GAGPRA/PSRAGPSVGGLGLRP
1		1		I		SGRAPRAEVALPAGLSPAGLAG
1		1		1	l	PPPSGRSTGALPPAAAPTTSGP/P
1		1		I	l	ASTPPNAPAPGHTLSSRSPHRA
	l	1			l	ARPPPHCGAEEWTGFTMRPSW
L		L_	L			RISQSFIVKDS
14258	44626	Α	14341	482	700	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
_		<u> </u>				
14259	44627	A	14342	1368	2416	SSMTQSSCPMTSYLKSKFGEPL
		i				MPSYTNPTETGVSKLVNKTQVF
						TYAHKGPVSIYFDACQAAHLSK
			ŀ			LNNIWT\SVKI*DKKESASRATK
						AITEESKKECPDCDNQWTTHEF
						NQHLYTGRAALFASQEEKIGYT
						TGTCYPLNLTILKPNMTFWTKG
						HKGLLTFDQAGALLGLGIPLVI
						TKKTQRTQVQLSPIQQFRFYKS
						FNEHFNSEVSKIQIPPISTENLFV
				ľ		QLAKSIANNLGVTSCYVCGGA
			l			NMRDQWPWEARELMLQDNFT
						LPEFVTKFNANPSVWLLRNPIIG
						KYCIACWGKSFQNQIGETTCLG
						QQYFEESENRTQWRSFIDDSS\G
						HFNPFLQFPTLNQSWYQLE
14260	44628	В	14343	I	1071	
14261	44629	С	14344	89	283	
14262	44630	Α	14345	244	315	NPKGQKDREAPLSRSVFVLKIKI
				l		KRAFALLLHGRF*PSTPAASASP
						ASKSPRGSGKALASALFYIVQL
						TK
14263	44631	Α	14346	1228	1656	QVYRPSQTPHLALSPERVAPGR
				l		RAAGRLAPEARAPRGSP/LPPHR
						VSEKTIRVVVFHPGARRAGGTP
		l				PRAPRGDTGGAPGA\PTYSTPL
			l			MSLHRARLESSPTGSSFPADSA
		1		İ		KPVPLAVVSLDSR*GQWESRSS
		ŀ				IHA\VTN*MTRH
14264	44632	Α	14347	2	705	GSWPGLDDGSATRILACPWPGL
1		l				GDGPAAPKANLPLVRWKPQCF
		1				RGEGL*GSSCFSATLLTGLDLW
		l				ACCSSRSTASARLSTYPSAFTSG
		l				WCSSLGPTEEAPGSWINCPLAG
		l				ILGAATSPKPSAIPQASPEVSLD
1		1				AEDTGMKP*DSA\GPPGPARGL
		1	l			CGRGCGLSPACMGLRP/PSPPA
1		1	l			QSPSTALTSLT\PSPFHPPRKKW
		l				SPPSAPAPSAPSPAPASLTPSPAP
1	1	ı	I	I		PAPSPAPTPAPAP

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop cudon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14265	44633	A	14348	367	1174	WVRGQPSQSQLDLLACSTPTS*
		1		i		WPPGRAPPPPPLP/PPPPLPP\PTP
		1		ĺ		ASPPLLHPPRRAPPPPPVP\SFTP
ŀ		l		1		ASPPPDPVPRPTKALTGLP/LCP
		1		1		LPCMGR/PFVPYLGGLLPCLGP
				1		DFSSL**SPGGWDRPVRPLGISQ
		1				IPE\PPLTMGL*MQ*IPP*PRFQG
						HYAPLPPLQTLCISVKTPGRNPR
	1					GLPMHKILQPQNPSQPPCELPQ
ļ						AHCPTTPPPKQCP\PAAFSILPVT
						FADKDPGAAPEGTVKAPMPTQ
						*EL*PLRPREHRVLSGPTAGQT
14266	44634	A	14349	332	988	RRAGTPASWGHNGARAKTPSR
		1				ACACLGLYPIHQPTARRAPAAR
	l					SHGPGPPALPRKGAAHGPRSVS
	İ	1				HHPLRPCSRSSCLAWPCLHHPQ
ŀ	l					PLHCHPHRHWPKWHPHPLAPH
	1					PLLGLHPRSRPQTPPWTSCGPSR
	ł					RLPTPSGSWPAPFDRDWPN*AR
		ł				PSALCCPFCQEPQLTPCLHLCPH
						PHPHRHLPGLSCPHRPPRWRSP
		ļ				QSPCPWWLLWWTGQWWQPGE
14267	44635	Α	14350	190	718	FVLQLPPAPKSGLGRWGSPPPG
	İ					QFSSSQRTGTVRGRGRCAEPRG
		1		i		SLPHPT*GNNK*HGNWMGLGS
		1				QGL\GRQPGEDKHCYQR/IGVG
	1					GAHLLRPPHQRIRSLTSTGSGCS
	ì	1		i		HPWSRELGCSWDGQRASAPLA
			1			LRQAWGEGLAGRNHTIVPFHG
		1	i			DISIAPRLLPCRETLFSALCGGES
14268	44636	Α	14351	1	486	MKGEVSCKQHQDALQQVLWA
	1					MLQDADKQVRRAPGPVENENH
	1					CDFVKLREMLICTNMEDLREQT
1	1	1			1	HTRHYELYRRCKLEEMGFTDV
1		l				GPENKPVRVLLFKSLSDSKRVK
1	1	l				YDEEWIRLPVE/MKY/MLERKV
1		1				TQKQP*VPLVFVSWRSR*ESLR
1	1	l			1	REEEENQQCHVLEL

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	,	
14269	44637	Α	14352	1	2380	MMWSVLLQPVVLAAVERQGE
						ALEFLIGVPESDGENGTKLENT
						LQDIIQEDFPNLARQAHVQIQEI
						QRTPQRYSLRRATPRHIIVRFTK
1						VEMKEKMLRAAREKGRVTLK
ł						GKPIRLTADLSAETLQARRDRG
		1				PIFNILKEKNFQPRISYPAKLSFT
		1				SEGEIKYFTDKQMLRDFVTTRP
		1				ALKELLKEALNMERNNRNFFG
1		l				VTFAFFVGGQSLGKALQETSIPF
1		1				LREEDLEKTSILETGNGDWEAT
1						RGLPEECAGMRERDCQQNEKY
1				1	1	QVPQFDQSTIKNIESAKGLDVW
1						DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
		1				KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
		1				LTTAQVNVSKSDDTLKINGVEL
1		ŀ				HKTIFDGDGKTYQNVQQFIDEC
		1				NYTSGDNHTLRDPHYVEDKGH
		ŀ				KYLGLIPTYDISSNIYLISKSCLW
		ŀ				QANFRLNSALAQSPAITNFQFR
		ļ.				GHSLKTLDLLTMKNLDSKVNII
ŀ	1					PVIAKADTVSKTELHKFKIKFM
		ŀ				SELVSNGVQIYQFPTDDDTIAK
						VNAAMNGQLPFAVVGSMDEIK
						VGNKMVKARQYPWGVVQVEN ENHCDFVKLREMFICTNMEDL
						REQTHTRHYELYR\RLQLEEMG
		1				FTD\VAPKNKPASVQETYEAKR
						HEFHGERORKEEEMKOMFVOR
14270	44638	A	14353	105	328	TIET HOERQIGEEEEMRQIMI TQI
14271	44639	Ā	14354	107	676	IRPDSLPSASPAQDFYAASSNTF
						PA/DRWSALTKIYEDQTSHNIPL
						SQ/TLIQSKALTLFNSVKAEKDE
						EAAEEKFEVTRVVHIRGLIDGV
		ŀ				VEADIVEALOEFGLISYVMVMS
		ŀ				KKRQALVEFEDVLEACNAVNY
						TADNQIHIAGHPAFVNYSTSQK
1						SRPGNSDDPEREQCASLYHPEP
		ŀ			1	HLFDHHGCSLHYR
14272	44640	Α	14355	221	648	LDWLRRVVGIPSLRWTCATLV
					1	GTRRAARHRCVNVWECGGLKE
		1				ITY*SVEGLCRA/EKPKPLALLE
			1		1	ETCKLPERQVVNAKEKFLEE\IK
		1	l		1	SATPVNT*MIRKRNCIITDMEK
		1			1	V*AVWIEDQTSHNLPLSQSLIES
		L				RALTLFNYVMA
14273	44641	Α	14356	122	317	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nocleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14274	44642	А	14357	338	489	WLGQN*RHGKQPGPGVGHVGT TVWS*PHSPACWAPAPQGPAPT GEEGCHF
14275	44643	A	14358	1	459	RPVWRNSQPPTAAYRWASGSR MWSTSFSRWNKTPSGKACSSS DKGPPMAQVPIPQRNVPPTK/ QPLVLCHGEAPDFSRPPQHRPS LQKHPSTGG/PSHSPLPTSAPSSR AGLPPLAMTLDLTSGWRLWTPI QPFRLPPGSCREGTILLCSVMD
14276	44644	A	14359	609	2077	GLKAQPSSSRGHSRGPGACAAA PASRVHLBVTFCPEPNSSPCPHC TGGWGGRGQ/PACAGNPTASSP CQIECPALRSCLHALMDLGSPQ/ LRGAPRTGVPOPRLHLLPQPAL SSPPVLLRGPHLRGTCPLAARQ GQSPPAPGGLRGHPGSHRGRLPP AQGGHRLPQPPGGRRHRAPQP DPRRFGLRAAAASGTGLAAWG AVRAARERSGGLPGGAGPHH ARAQPGPRAAPPAAAAAFGVAP AGGGAVPGGHPHRPGGEAALP HGLPAPAAERTAPRIQPCPAC GLIARSDVHHPPVLPPILSGPAP ACRLVGQLGVRGAPSSPVPHP ALHFPRAQP/PPGKGITTTSQTQ PAGSPSPCMPPAATPGARHPP AGGNGHGLGVGGAGFHGGHHGGHHGHGHHGAQLL*DQAAVGVRGP GPWALSGMSQPTLPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPHP PACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGHPAP ACGVRAFQGTBPTAGTPTAGTPTAGTPTAP PAGGPWALSGMSQPTLPTAGTPTAP ACGVRAFQGTBPTAGTPTAGTPTAP CGVRAFQGTBPTAGTPTAGTPTAP CGVRAFQGTBPTAGTPTAGTPTAP CGVRAFQGTBPTAGTPTAGTPTAP CGVRAFQGTBPTAGTPTAGTPTAP CGVRAFQGTBPTAGTPTAGTPTAGTPTAGTPTAGTPTAGTPTAGTPT

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14277	44645	A	14363	47	1250	EPRPSRSSGGPQLGGGLPRPMK KIPGSSAKTAGPLATREPVPRAP *SAGRQPWFQRPWGKRKGRKT *NHGSPGVKPTPGPLNKDKGE KEERPRQQDPQRKALLHMFSG KPPERPLPNGKGSTESSDYLRA SYTPGPWSFSPQVASGPMPV TTKKPA*APPQVLEPKERQTGA AADMQPQAWHQGREPLLV PTHSRPEGGCREVPQAA/SQNPR PAPGLQTPGTRGTSSVTPQCP PAPGLQTPGTRQTSSVTPQCP PAPGLQTPGTRQTSSVTPQCP ROPPHSRPCLFTAQACTMSHHS RQPPHSRPCLFTAQACTMSHHS RQPHSRPCLFTAQACTMSHHS LELQPSTSPQMGRTTAQACTMSHHS AAGHDG/DPASQSALPETGKRT LELQPGGPLISLS*EAGSLSRSE PSCVREV
14278	44646	A	14364	23	714	EAPEIPRQNRRMTMKERPVWL PHIK*IHPNVAKWSRVPVQVQK NPQPQBVPTPEFIPVPGTSPEAAG FWH*GAHDSFGTRDRSEVEPGR GHPHLGPRDSAAADPGEHSPCD CP*ACPEVRRWPA\APVQIQYIIP SVDDFSLEFHAQDGDISDMRRE NVPSSPAEGKAAPLYQQPLMI PQANIMAGISPS\SPTAAEFQHR DVHHRGPGTRDRTHTETGRVC PHPVPTDSPAL

1782

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14279	44647	A	14365	3	1869	ITTVAPSRPASSDCESDGS*EOD
14277	1707/	ľ^	14303	ľ	1.005	AGPGSRMGPEK*AGPRPTRRVG
						RWSQTHSQTDDTSGAVVPEYP
1		l				SDR*YAGGRGPG/PGGPTVHP*
1						KRRS*NPQETKMMARRDPKSW
1						
						AKRLSPV*PRLAALLTSASPSCP
						LEPRVLPFRVPPSHGLCEGTSYP
						NSPGT*RPPCRSSAKTAGPLATR
1						PEVPGAP*SAGRQPWFQRPWG
	1		ļ.			KRKG\GKPETMEAPG*SQPGAL
	l		İ			EQG*GREGRETKATRPAEEGSP
						PHVFRETSREAAAEWKRIHGVF
		1				*SSEGHPG*FPFSFPSAGGNRGT
						PLSCLLGVRDSTILPGCKRANA
	ĺ		İ			GPHNQ*EAARGPCPR*S\SAAE
						MSGRGSVLASLSPLRKASLSSSS
						SLGPKERQTGAAA/AHPSACSQ
				ł		APGPRASPRGEADTQPPRG*LP
				1		RSSPGCLQNPRPAPGRQTPGTR
		l				QTSCGDL/SSPARQPPHTAWA*
		1				APISASGQEPRDLP\SSDSGLPEL
						PQETETGSLPDPRKRHPGR*AG
						GPGKSPTSASRNRTWTKYVAP
						DGQEDTGPGAQRRPAASAQQT
1						LPAYCPGLHHVPSLSGQP*WGP
						ASQS/GSSGDWKTDAGAPASW
						RPPHFTLLRSREPSSLRALMCQR
			ł			SLRLPVFVSHRASSMRTFRFPPP
		l				QRTAILTWS
14280	44648	В	14366	1	1519	
14281	44649	Α	14367	312	1326	TTKTGQDKDRLEEFSMQQLSGS
		l		1		KRKMSTQPRTEL\PPNRHTTTTS
	1	l	1	1		TTPATNTTCTATVPPQPQYSYH
	1	l		1		DINVYSLAGLAPHITLNPTAHP
	1	l		1		QLKQCVRQAIERAVQELVHPV
	1	l	İ	1		VDRSIKIAMTTCEQIVRKDFAL
	l	l				DSEESRMRIAAHHMMRNLTAG
1	I	l				MAMITCREPLLMSISTNLKNSF
	ļ.	l		l		ASALR/PDGSSAIQL*SND*SCS
	l	l		l		WRPKLYDAFWDLSSLRV**PSR
	1	l		l		PEGEGRVSSEGMGESLPFSSSW
	l	l				PROYOSFLCICWTDAPARNTED
		l	ĺ	1		R*SHNKVLSSVY*NVC*NOLPC
		l		1		SG*AAAQSCCQSHHDPSQVLSQ
				l		PGCLCSTHCTARETLRGGHQHC
				l		HKD*SAEQGL
					L	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first eodon for peptide sequenee	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14282	44650	A	14368	2	1120	PEPEPGARGGEQLNEDFEG*VA VPTCLHCPALSKTQSRAIQALT VRGEATIRPPCDDTKGGHRF LNELIKVVSPKVGAPSPSSDLPC RWWGEERGWTVGLPEEVKLEA AYQMLKKQGEAHRGWGATRA CLPLPHHERRKNVIFEDEEKSK VRLQGGTVGDSEPRWGSVGSI KEMVQFDSGRAQSTGRRDG GIELGHNVKLLTEMVMSHSQG GAAAGSSEDLMKVHLPPCLPHS LPTPOPTRTLRMEPTLFRLASDT EDDDEALGEPRAGGEVRSPPLP PPPAPVRGEEVNGDATAGSIFG EEVAGELGGHPSGWRGTESYP AMPTRRGEQASPQPSGWRGTESYP AMPTRRGEQASPQPSGWRGTESYP AMPTRRGEQASPQPSSWSLL
14283	44651	A	14369	1	432	DDELMSLGEEGARPGGGRAQQ GTRQEVTAGVEPAREWQRRLM EVQWGLERAPYCIQHAPR*RRS PARPGTQSCGLGACWIQ*GALS SPHCTSISRRCHSLAGSTPAVTS CLV
14284	44652	A	14370	16	471	
14285	44653	В	14371	66	206	
14286	44654	В	14372	234	390	
14287	44655	В	14373	207	427	
14288	44656	А	14374	2	572	ELLERAPYGVEFSLLPGLWLGS RAGVGRHDKRYGLGRAPTWS LEGAYSSMTPRPOLCVP/WPC WGPSSPGG*QQLGQISCSYSLV DR*VALLWFWMLVGSPLVKGH CPTPWSRRVDFSVPLATPGILYY WGLFAVFLFSHLLPARPGEOMS ACCRHLHARSPLLTIDSLIKYY SCNSQGKLVRVFPPGD
14289	44657	Α	14375	175	499	KQNKSMGRKSQGFVASFGNHL YCTASCGMPASSTSTQTWPRVL PLHSQ/ESSHGGCSWRGRR*GP RPRPHKTPVPMVMDSTARNRA RRELVLAWCKSPLATALLVSW DK
14290	44658	A	14376	408	625	VFGEQEVFGYMNKFSSGDF*DF GAPTTLSSEHCTQCVVFYPSAP SFPFPPVPKVHCVIPHSLVPTYE WEHTM

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14291	44659	A	14377	1263	1812	AGDAAPEDPAGRGOEDLYDEE
14291	44639	^	14377	1203	1012	
1		ı				LPGWGLCIPHPACLQPCHA*VT
		1				KAG/ARVKVILPS*DTQRTKDIL
		ł				KAGGG/PSSIHFINSAHGKPAVI
						VRPRGIWVLNTTPLSGQWGPYI
				1		GVYTAATVDMKSLGSGSLGLQ
		1				LALPLAASWLGVLTEAARVFG
				1		EQVVFGYMNSFFSGDFWDFGA
						PITRAVYTVPNV
14292	44660	Α	14378	1	693	
14293	4466I	Α	14379	2	1299	WDYV/KRPNLRLIGVTESDGEN
1				1		GTKLENTLQDIILENFPNLARQA
1				İ		NIQIQEIQRTPQRYSSRRATPRHI
						IVTFAKVEMNEKMLRAAREKG
		ı		1		RVTHKGKPIRLTADLSAETLQA
		l	ŀ			RREWGPIFNILKEKNFQPRISYP
		l		1		AKLSFISEGEIKSFTDKQMLRDF
		l	l			VTTRPALKELLKEALNMERKN
ı		l	l			RQINETENQQGYPGIELGSAPSG
1		l				PNPLFCRHLQMSVDGRHLQNSP
1		l	l			POINRIYIFLAPHHTYSKIDHILG
1		l	l	i		SKTLLKKCKRTEIITNYLSDHSA
		l				IKLELRIKKLTONCSTTWKLNH
	1	l				LLLNDYWVHNEMKAEMKMFF
						ETNENKDTTYQNLWDTFKAVY
		l				RGKFLALNAHKRKQERSKMDT
1		l				LTSQLKELEKQEQTHSKASRRQ
1		1			l	EITKIKEELKEIETQKTLQEINES
		l		1		KSWFFERINKIDRPLARLTKKK
						REKNQIDAIKN
14294	44662	Α	14380	2	383	REKNQIDAIKI
14295	44663	A	14381	1	1536	
14296	44664	A	14382	2	1118	REHLAEGMAVGTASADLNISA
1	1.00	ļ.,		-		CWLRREQQISQHSARALLRDRL
		1				PPQENSSWHPAGAPLGRNFQRK
		l				EOAAIFAVLOPPLVIPRKTGSGV
		l	ŀ	i		DLEOTPADLOKRGLTVTRKTN
		l				
1		l				KQKAIASTSTERTPTQKPPFKSH
		l				QHQRPNVDKSAKMRKNQSKK
		l	l			AENSKNQNTSSPPKDHNSSPAR
1		I	l	1		EQNWTENEFDELTEVGFRRWV
1		I	l	1		ITNSSKLKEHVLTQCKETKKLD
1	1	1	l	1		KRLEELLTIITSLEKNINDLMEL
1		1	l	1		KNTAGELRDAYTSINS*IDQAEE
1	1	1		1		RISEIGDQLNEIKCEDKMREK\R
1		1	1	1		MKRNEQSLQEIWDYVKRPNLR
1		1	l			LIGVPESDGENGTKLENTLQDII\
1	1	l	l	1		QENFPNLARQANIQIQEIQRTPQ
1		1				RYSSRRATPRHIIVRFTKV
$\overline{}$		•				

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
14297	44665	IA	14383	1	1154	MLPILGSRHPSSLGQARKLDYR
14297	44003	^	14303	'	1134	PSKSOCLEPGTPRACLVLYTLV
					i	AKLIYNDYFIHYFAPRGLPPME
		1				KNVVFVIDISGFMFGTKMKQD
		1				HNFSLAREQNWTENEFDKLME
					1	VDFRRWVITNSSELKEHVLTQC
						KEANNLEKRLDELLTRITSVEK
						NINDLMELKNTAGELREAYTSF
						NSCNDQAEEKISEIEDQLNEIKP
		ļ.				NLRLIGVPESDGENGTKLENTL
						QDIIQENFSNLAKQAN\VQIQEI\
						QRTPQRY\SSR\RATPRTH/IIVRF
l		ı				TKVGNGREKMFKGQPGGKGR\
		1			i	VTLKGKPITLT\ADLSAETLQAR
l		1			I	REWGPIFNILKEKNFQPRISYPA
l						KLSFISEGEIKYFTDKQMLRDF\
				1		VTTRPALQELLKEALNMERNN
l		1				RYQPLQNHAKM
14298	44666	В	14384	1	1554	
14299	44667	Α	14385	1	909	
14300	44668	Α	14386	2	207	DIMSCILQNYNRPPVMALAIPIA
						VKFLHRGNKELCRNMSNYLSL
						AAITKAD\LLADHTEVIVKSILQ
14301	44669	Α	14387	496	1207	PLITGTHDSGSVESQLIKLILTM
					1	HQLFRLVLGQKDLSRAGDLF\S
		1				*NNSEIEDSLTEALEQIKIISSSSD
						YQTNNNDQAVVEICITRITTAIR
						ETESIEKHAKALVGLWDSCLEH
	l					NLRPFGKDEDTPHAKIASDIMS
	1				1	CILPELQPTPQSMALAIPIAVKF
					1	LHRGNKELCRNMSNYLSLAAIT
						KADLLADHTEVIVKSILQGMVR
		l				KLSLGTCFGRYLKVFSSSIYGL
						WEARPRVLEAN
14302	44670	A	14388	219	448	VSLELCLKTTGPDCGCQTSQTP
117502	11070	1	1.500	[PAAAYHHL/PGAGLPCGPRLPA
	Į.	1				LCCSCPMGARPL*HHKR/DPSRF
						QEKTHLYLPSSH
14303	44671	A	14389	542	1365	GTEGVMANRSQEGNGGGSWR
14303	44071	^	14309	342	1303	AKOVVVTVPF*EQSLVIMLWR
		1				RGGNT*ORPSEGTSRSHICSFH
						MSRHCQ*CKCLLKEP/VSAEAR
						DALDTKTHFPLAMIPSYLLLNG
		1	Ī			NIGELPEGPAGGCAQNPGLWAS
		1	1			/RGPARR*RSQASSSPEVRWPPG
	1	1				QCGLRGRPCKKE\PRPHPWSRP
						CASPYRGAGGQRLRRACSAAG
		1	1			RKTPDRRPLPEPTRGPVSQRAV
		1				LGKLRAAAGHRIPIGDLGDGLS
						LGAPPAGRHLHTVLPLRREFSP
		L		l		CEDALGIGEQQGKKRVT
L	Ь		L			CEDALGIGLQQGKKKYI

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14304	44672	Α	14390	3	723	PVLRGGIQEFAYIKALYERKFL
						VPKPIDYNRHAVDMELINGYPL
		l				C*IHHVEDPASVYDEAMELIVE
						LANHGLIHGDFNEFNLILDESD
		1				HITMIDFPQMDSTSHPNAEWYF
						DRDVKCIIDFFMKRFSYESELFP
						TFKDIRREDTLDVEVSASGYTK
						EMQADDELLHPLGPDDKNIETK
		ŀ				EGSEFSFSDGEVAEKAEVYGSE
						NESERNCLEESEGCY\SDHLETL
i						NK*RKTVYQKRVLMHGVLK
14305	44673	Α	14391	I	1889	MGSKVKEVEDDSQMSDLVRG
		l				MNAFSKTGSGGVEQGRTGQGG
		l				TNGHIKKAKARDRDRRFKKTP
		l				QRKVGNDNLQLPLRWTEQHAE
l		l				THIMNFCSKNYNRNIPGKPRESI
1		l				DTLNEAACRGRLSKTARKLWS
		1				EAQDQRSFQATEMQAFLFSAK
						RIAAEKKTIIRRRDLCSIWGKLL
		1				PRPGSQSVPAQIPENLQEQEERA
		1				KWLLNGRSKFFPQTTSTLGKVT
						SNKTGVKEPELESRQPNLLKVE
		1				MGMKNHEIVPGSLIASIASLKH
		l				GGCNKVLRELVKHKLIAWERT
		l				KTVQGYRLTNAGYDYLALKTL
		l				SSRQVVESVG\NQMGVGK\ESYI
		1				YIVANEEGQQFALKLHRLGRTS
1		1			i	FRNLKNKRDYHKHRHNVSWL
		1				YLSRLSAMKEFAYMKFRTLEIR
		1				DYFYWVKINVLTGPCSPKRGSS
		1				GEFVPCLLQLLLGFGDCWHSFG
		1				LWHITPVSVSMLTLLPSLLLVK
						QKKPTLDSRMGQFDFVVELQL
1						YLCVLEREVIREYPPDGQIFGRY
1		1			l	FDRDVKCIKDFFMKRFSYESEL
		l			1	FPTFKDIRREDTLDVEVSASGY
						TKEMQADDELLHPLGPDDKNIE
						TKEGSEFSFSDGEVPEKAEVYG
l	1	l			ŀ	SENESERNCLEESEGCY\SDHLE
1		l				TLNK*RKTVYQKRVLMHGVLK

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14306	44674	Α	14392	150	1910	STPPGGPSVLM*STEEPGAGRS
			'			QPLCDDLSTRPLSCLGAGCCLW
						PAKRISAKEEPWYSSRSKDKKE
						KN\YSSPETTTSGGCHSPGDSQY
		l				QELAVALESSSVTINQLNENIES
		l				LKQQKKQVEHQLEEAKKTNNE
		1				IHKAQMEQLETINILTLEKADL
		1		İ		KTTLYHTKRAARHFEEESKDLA
		1				GRLQYSLQRIQELERALCAVST
		1				QQQEEDRGHCLSSPDQNFSLFT
		1				QSSSCREAVLHRRLQQTIKERA
		1				LLNAHVTQVTESLKQVQLERD
						EYAKHIKGERARWQERMWKM
			ľ			SVEARTLKEEKKRDIHRIQELER
		l				SLSELKNQMAEPPSLAPPAVTS
					1	VVEQLQDEAKHLRQEVEGLEG
		l				KLQSQVENNQALSLLSKEQKQ
		l				RLQEQEEMLREQEAQRVREQE
						RLCEQNERLREQQKMLQEQGE
		l			ŀ	RLRKQEQRLRKQEERLRKEEER
						L/TKAGKEAVGPGGEAVEEGGE
		l			1	ATKAGGEARALPEPQARQAAG
		1			ļ	RATVQLRGSEQREKERTAVGA
		1				ASKGAAGEARRGEGDAVHGHL
		1				SAADL*EGGAAQAVTAADPVR
					İ	GPAAAAGSLGQSGAPRSCQPPE
						PTARDPAKPRGSPWRRRWRTT
14307	44675	В	14393	463	3755	
14308	44676	В	14394	175	1835	
14309	44677	A	14395	102	465	RAYCWAYIRLSCLTVILIAWCS
						QESALSEEEEDTTRPLETVTFKE
						VAVDLTQEEWEQMKPAQRNL
		1				YRDVMLENYSNLVTVGCQVTK
						PDVI\SSWSKKRSPG*WRKKCL
		ļ.,		<u> </u>		GGTVQNPEEERTAV
14310	44678	A A	14396	1	531	MI I MIDOUTED OCDE A KNIMI A O
14311	44679	A	14397	1	231	MLLNIPQHTRQSPRAKNYLAQ
						NVNSVKAWCSQESALSEEED
		ŀ				TTRPLVRLKHLFVSCIILQCSYV
						LLPLTPEYQLNKFLYVVITMPN
		l		ļ		FDDFKLISIMFLQETVTFKDVA
		1	1		1	VDLTQEEWEQMKPAQRNLYR
		1				DVMLENYSNLVTVGCQVTKPD
	L	$_{\perp}$			l	VI\SSWSKKRSPG*WRKKCLGG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14312	44680	Α	14398	1	539	MPTTTTREMTQNSSETIKNKNG
		l			\	KKQRKQRKNSKKNRRKQKDE
						DKTRRQNKRHHKKNDINPEEE
		ı			1	RTAVRLKDGGAPAPQAVQEHD
						AGICSASGGAQETYILVEGKGE
					-	ASTSPGQSKGSREGGVGANETC
1		l				SKKLYRDVMLENYSNLVT/VG
						CQVTKPDVYSSWSKKRSPG*W
		1				RKKFWEALSR
14313	44681	Α	14399	322	1227	FDLKMSKCRKTPLQQLASPASF
1		l				SPDILADIFELFAKNFSYW\KPL
		l				NNEWQLPDPSEIFTCDHTEFNA
		1				FPWI*RTP*NEVKKPTGVIRTW
		1				DEWHEHTCFH**SGE/ISFPHGR
İ		ļ				KSVNAELWTQAWCKFHEILCS
						FPLIPQEAFONGKLNSLHLCEAP
i		1			i	GAFIASLNHYLKSHRFPCHWSW
				İ		VANTLNPYHEANDDLMMIMD
		1				DRLIANTLHWWYFGPDNTGDI
1		1				MTLKFLTGLONFISSMGTVHLV
	1	1				TANGSFDCQ*NPAGDQRMVKG
		1				HCYFPSWNWLHLSKHTQCGNE
		l]		FIFMTKKKOSLTAKSLPDTFS
14314	44682	Α	14400	3	183	KRKEITNSLDAAASRGTVPAAL
					1	AGVOYLEPRAGAAGRLGGAG
1		1				WGRVPRPGSS/SSLSLVGR*GW
		1				CPVLRTQGRCCRPPGGCWVGA
						RPKAWELLSLSLVGR
14315	44683	A	14401	21	399	IWNSAPTASRRPWSLAPGSPRS
						TWCVASTPPCAPGKAREPRDRS
						EVPPATAPAVLOSHWPCPRSSR
						SGSAGNCVGPPHSWAPGHSQH
						LTPGPGHGGPRGP\CWHLLMOT
	l					AWPALTPPGAMPGEAEGO
14316	44684	Α	14402	59	478	RCSQPWHCPAALAGVQYLEPR
				1		AGAAGRLGGAGWGRVPRPREL
						LSLSLVGR*GWRPQEQDQLFPH
						LRRPVGTRSESTWGNRAGAAT
1						W/RAATPSPWRT*NGTKSKQW
	ŀ					TASGLWQKGDGWD*GSGKPPL
1						PLSVPAGQRLVPVC
14317	44685	l _A	14403	1150	1319	PRRLPTACTP*RS/CSRYWMVP
1	1	1		1	'	KSPRMTMMMWTKLARMGAQ
ł		1			1	W*PRKSNACLSRAASS
14318	44686	Α	14404	34	304	LPSRGAGLGTCSPPCLSLPPTPW
1.7510	1.7000	ľ.			I	APVRPEPPRRAPPPAPRRPVPST
	1	1				TO/EARRSAASLLSPARPRAHRD
		1				GTNNSRRATLRAVTLTARVRG
1		1			1	F
		Ь.	L	L	L	<u> </u>

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop coden, /-possible nucleotide deletion, \-possible nucleotide insertion)
14319	44687	A	14405	1	360	SSSLSYLPAGQGSGPAARHA*A SHPLHGLLCGPSLPNEHRPLLQ GAQSH*PPKG*GVRAHGVGLA GRSTCSP/ECGIH*VKPAGLLSFL GPRCFYEL*QSPRRSAAASLLSP ARSRAHRE
14320	44688	В	14406	80	1166	
14321	44689	С	14407	1	543	
14322	44690	В	14408	1	930	
14323	44691	В	14409	1	1473	
14324	44692	Α	14410	388	606	
14325	44693	Α	14411	1	798	
14326	44694	A	14412	687	832	
14327	44695	В	14413	1	1218	
14328	44696	В	14414	1	642	
14329	44697	В	14415	1	807	-
14330	44698	Α	14416	554	724	
14331	44699	Α	14417	216	494	
14332	44700	Α	14418	1	3408	
14333	44701	Α	14419	330	858	VPSLDGQRGLELDTECRLVYLQ SLS*T*RFSTSSPEQLDTECRLV HSQTLC*TQGADWCIYKP*VGL QVELPASPAP\WLAFLSPWVVN GTGRHGAGGGARRGGSGCTGA HGVGGRLRHGRLQVPSPAPWE GSCSLPSPIGPPFKRAHSVLQKG SSTLMVGTCASSPKVVPFHLLS
14334	44702	В	14420	182	1513	
14335	44703	В	14421	1	1608	
14336	44704 44705	A	14422 14423	167	1830 433	LPSRGAGLGTCSPPCLSLPPTPW APVRSEPPRRAPPP/GSTVPSPIN HPRAEECERMARDWQAAPPAA PVRDPLGEASWAPEFFGSMLPL
14338	44706	В	14424	32	352	
14339	44707	В	14425	103	451	
14340	44708	А	14426	158	921	LIPSR WAGLGTCSPPCLSLPTF WAPVRPEPPRRAPPSAPRPVP, RPPKG*GVPAHGAGLAGSSTCS PDICRVCRSEGTPEKPLYHPCVC TGSIKFHQECYSPDMPSRLPIQ DIFAGLVTSIGTAIRVWFHYTLV AFAWLGVVPLTAWHQLLDLGS TLNPGCSHLVILKLTTSTKTLFP NKATFILVVLSQDKTLMEDPR MEDKWKEDRFSPFNSEALEITL GERHRPQTASVIPWFCSSRHVL NLGKINFEVD

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14341	44709	A	14427	242	515	I IQSVHWCIHKP*ARHRVLIGVFT
14341	44709	l^_	14427	242	313	IPELDIKVLHVPTRLRNPAGFTH
İ						WILHRDCRCSCMPVLRHAPALL
						SPWVVNGTGRSRAGDSTHRGG
		ŀ				SG
14342	44710	A	14428	759	988	TWPRKPLFPPRPLGL*WEGMLO
14342	77710	n	17720	/32		RSLTCPGDIFPIILVISIWLLVTY
		l				ANFGSWL*FLPRKWVFPFFIPVP
		1				NKFLIFI
14343	44711	A	14429	130	788	VLQLIKAVWTPRTQEPSWLPPV
14545		ľ	1.7727	150	1,00	NPARELQVELPASPAPCARTPQ
1		1				LLGGRWDWAPWSRGWRSFGE
		1				ARAAOKPMEGVGGSGMAGCR
		1			}	SRALPHGKAAKARRAIEHSTGG
						LALLGDPVHSPQPLARVLSPSLP
		1				RASRAGWLLRVRGPPSPRPPGT
		1			ŀ	PAGRQAPHAAPVPARASPSTPP/
						CRAERVGSSPGQPRKGLPQCRG
l				İ		EAEGLLKCHQSGSPGRGEELW
						HSR
14344	44712	A	14430	108	1686	ANRCPLPHPGTGAGRCLGASA
		1				GRAEEAGV*GPLHEGLGYRPA
1		i				GLFLRADTGHRTPGWGGGGGG
		i				AGGRGGAAPGPGVGATRRFAG
	1	l				RRGCARHGAAVPAAVR\CCERL
į.		l		l		VLNVAGLRFETRARTLGRFPDT
	1	1				LLGDPARRGRFYDDARREYFFD
1		l		l .		RHRPSFDAVLYYYQSGGRLRRP
l		1				AHVPLDVFLEEVAFYGLGAAA
				l		LARLREDEGCPVPPERPLPRRAF
		l		i		ARQLWLLFEFPESSQAARVLAV
		l				VSVLVILVSIVVFCLETLPDFRD
l		l		l		DRDGTGLAAAAAAGPFPAPLN
İ		l				GSSQMPGNPPRLPFNDPFFVVE
		1			ĺ	TLCICWFSFELLVRLLVCPSKAI
		l				FFKNVMNLIDFVAILPYFVALG
		l		1	1	TELARQRGVGQQAMSLAILRVI
		1				RLVRVFRIFKLSRHSKGLQILGQ
				l		TLRASMRELGLLIFFLFIGVVLF
	1					SSAVYFAEVDRVDSHFTSIPESF
				ŀ		WWAVVTMTTVGYGDMAPVT
	1	1			l	VGGKIVGSLCAIAGVLTISLPVP
						VIVSNFSYFYHRETEGEEAGMF
						SHVDMQPCGPLEGKANGGLVD
14345	44713	Α	14431	1	427	
14346	44714	В	14432	8	323	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14347	44715	A	14433	20	1050	FNLVYMSHVSDEGALSESASAT
						AASWPPHGFLOHFVPGRPOVPI
ŀ		l				CLYLLGOKTG*HMAPLLROVT
l		l				GSPIPPKTPGRGIRNMKRNEKR
						RYKTLMADGWELHOELSLGRT
		l				SSLPPYKGPGPWSPGFQKVSKG
		1				ANPVEIKRGVMLAVDAVIAEL
		1				KKQSKPVTKPEEIAQVATISAN
		1	ĺ			GDKEIGNIISDAMKKCGRKGIIT
		1				KISSVOSIVTALEIANAYCKPLV
		1				IIAGDIDGEALTTLILNRLKVGL
		1				QVVAVKAPGFGDNRKNQLKDT
		1				VIATGGEVGEVTVIKDYAMLL
	1	1				KGKGNKSQIEKCVQEIIDQSDV
						TTSEYEKEKLS\EKLSDGVAVL
		İ				KSHPWSFCTKIYPLT
14348	44716	Α	14434	3	764	TTATVLARSIAKEGFEKISKGA
					l' .	NPVEIRRGVMLAVDAVIAELKK
		1				OSKPVTTPEEIAQVATISANGD
1					ļ	KEIGNIISDAMKKVGRKGVITV
		l			i	KDGKTLNDELEIIEGMKFDRG\
				l		YISHPFINTSKGOKCGISRDAYV
		1		ł		LLSEKKIS\SIQSIVPA\LEIAHAH
		1			i	RKPLVIIAEDVDGEALSTLVLN
				l		RLKVGLQVVAVKA\PGFGDNR
		ľ				KNOLKDMAIATGGAVFGEEGL
		l			i	TLNLEDVQPHDLGKVGEVIVTK
		l				DDAMLLKGKGDK
14349	44717	Α	14435	1	3288	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
		l				EAADLVGVSSQAIRDAEKAGR
1		l				LPHPDMEIRGRVEQRVGYTIEQI
		l				NHMRDVFGTRLRRAEDVFPPVI
1		l				GVAAHKGGVYKTSVSVHLAQ
						DLALKGLRVLLVEGNDPQGTA
		1	1	1	1	SMYHGWVPDLHIHAEDTLLPF
1		1			1	YLGEKDDVTYAIKPTCWPGLDI
1		l		1	1	IPSCLALHRIETELMGKFDEGKL
1		1	l		1	PTDPHLMLRLAIETVAHDYDVI
1		l				VIDSAPNLGIGTINVVC
14350	44718	A	14436	2	422	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=passible nucleotide deletion, \=possible nucleotide insertion)
14351	44719	A	14437	529	2484	WGINIIKKNKKAAPRAFGVRLE ECQPATENQRWQDLNVISSLLK SFFRKLPEPLFTDEELPLAPSFCP EEENGGRGGGPSPRYRLEPMDT IFVKNVKEDGPAHRAGLRTGD
						RLVKVNGESVIGKTYSQVIALI QNSDDTLELSIMPKDEDILQLPA ERESGGEASEPPRVVRPEPSTRA LEPPAEDRGDEVVLRQKPPTGR KVQLTPARQMNLGFGDESPEPE
						ASGRGERLGRK VAPLATTEDSL ASIPFIDICVRPCA WLHALLDW MAQQR VLDMQGQSIGMGV SW DRELEEKGSEKGYDEPTSPSIDL
						QAKHYPASAVVSSAMNSAPVL GTSPSSPTFTFTLGRHYSQDCST SAHLNIFAGIFDISNDHLQVSTL ESNFTRIHLTHYTGGNFPAWLA WDA VPFSSDLLNLGVHVLETG
						AGTPFLGSACSSDGIASQMNTA GLPQIRDLPGHYYETLKFLVGH LKTIADHSEKNKMEPRNL\ALV FGPTLVRTSEDNMTDMVTPHA
						*PLTRSWETLIQALQTGSSVTKR YKGKENPCGLTRSLRQCPTLST FLPNIWQDSAPWRPGVSGPVGD LKDSTTCSSAKSKGSWAPKKEP
14352	44720	В	14438	1	1473	YAREMLAISFISAVKPQAQEAA GDAGAGQQHRRRLGAGGAQS LGRGPQRRGHQERS
14352	44721	A	14438	5	694	VPDKWGWCLFMSRYLETAIRT AACAASPRYLNPGRCSAARGPS
						SQSPRPPWSGHKKRTASPGWG GQSAASPMPPSRSWPASVGGPE FRLSETPSRPFTLSGMK*GWQV HSWAPALFCSVMSFSPDTCCLC K*AHSQPCSPSPSTVILLRETSSN
						WDWQPEDLGAPKASASPPPSTT RITRLRSSELQLRRLNPSDSCHR GFCFSFRCLLTPEHVLPKTVLRI QPGCKVA
14354	44722	A	14440	303	757	MAWKVSLRGRKVAGRAPSQLR WASPPSSGQSVKSLLLPPARRL CPGPGVPARCHRRI*GLGPGRG WRGEWRDTNPWEPPQPRSPS/ G/PNCAWSDGCPRGAPRRPRAL *AGAARRAAPARI*VPWRGSAG
						GCSNRRLQVPAHE*TPTPFIRYL

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, /=possible nucleotide insertion)
14355	44723	A	14441		534	RHEDHSEPTFLISPSLLCFLLPSA PPLPRLTPNPAALLVPAVQRME PVLFLPAPGAAACRVWHPALPP PRQAAVSSSRPCYIPCVARSAPA GRVRAAGVPPAPPSSSPRLGRV GGRSRSPNVVFRVRGAAPG/PG L*ISPWTP/PLNEAS*RQRGCPAS FAVYGGQEWRPPQWPGPPRGP
14356	44724	В	14442	157	2143	
14357	44725	Α	14443	258	4089	OLITICLSRPETGGKFEVDTORP VCPPSRARAARASAAVAAAA TUTPPTMDSFDLALLQEWDLES LCYYEPDRNALRRKERERNQ ETQQDDGTFNSSYSLFSEPYKT NKGDELSNRIQNTLGRYDEMK DFLTDRTNQSHLVGVPKPGVPQ TPVNKIDEHFVADSRAQNQPSSI CSTTTSTPAAVPVQOSKRGTMG WQKAGHPPSDGQQRATQQGSL RTLLGBGVGRQQ/HSGQTSVQC GGGPSDNGATTGGG
14358	44726	В	14444	1	1326	
14359	44727	A	14445	453	710	SQHFGKPRQVDHLRPGV*DQP GQHGETPSLLKIQKLTGRGGAC LW/QSQLLRRLRQENRLNLGGG GCSEPGSCHCPPAWATERDSVS
14360	44728	В	14446	32	290	
14361	44729	Α	14447	244	426	
14362	44730	Α	14448	69	221	QWRFCGIERGERWGKD*EIGW LPCLCGKK*IWETF\ILFCTKKN SSALGSC
14363	44731	A	14449	27	220	AVILTAKICSFTPEPSKTTSPPGG TNNSRRATLRAVTLTVKVCSFT P*GSAASFLKSVRPRTHQ
14364	44732	Α	14450	115	732	EPETEUDPLSWNSSQLPVAALH LPOMSSVGAQOA(CNLCCLGDL VSPSSLPALERTGPPPWNRGPK GRGRLPSL/TVSQPSJLVIPPGTG KSEVNADASSRPA YCSNLGRYS GPGPPSLVIPPGTGKESEVNADRS GPAVCSNHFPSALP**RSSMRS LSLQQTSA WTCRHFHTSILKSR WRFPRLNSCLLCTCRPNTTWKP CLGACTL
14365	44733	A	14451	528	875	EHGYFQGTLGQKGCKSGKPPG GTLKTLLVTLREPGRLQSP*TSL WVSERLP*YP*GDTGENCPKLS FHLRNLGPS*AG*RERPPQGGL YLPPQAYRLMQGSAFWPLFGY APSPQR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14366	44734	A	14452	2788	3414	ACFSYIAFWRINTTSTANLSKES LTSTTSRSWYNNVINFCTDWL. FYKWLLKNFPIYFYMIFFROIT FHSMGSCAARASPTSTTFCSTA PSPIDHFRAEESSTRRYTGRQL HL.QPAKICSFTBEPSKTTSPFGG TNNSSALP*EL*HSPRRSAASL QFRTR*YPYRYQSQGPGSH*L PLPLPSSHIDHF*LFHLQHTFQS HALLP
14367	44735	A	14453	662	1338	VLQLIKAVWTQRPLGGRWDW AFWSRRWCSSGR.RPHRNOWS GWEAQAWGLQVPEPCPAGRQL RPGEKSSTAPVCTSCCRTRSTA GPTGCPSTRRDPTLSASPSTT ISLATATASASMWAGSCTSIST RGSASLVHQALTWPGSGPASPR KRYGSHPVA-AGVRWRAHRSL QPQTPGLK*SSHFSLPSGWNYR HSNHPISQSFPHLSPLSIPQNHH CHHGPFSW
14368	44736	А	14454	3	415	DFLLFSEILPASQSASAILPALPA SHRCPCCLDVPLPPEVPGPARLY VAAASVWRPVQHLSLLGRWA GPT*DRTESRETNAEVEADLPP LYAHIGRLEPTAGPSGASAVLST SSEYCPGPHPKMGHRRVTSCLS TCQS
14369	44737	Α	14455	659	818	
14370	44738	Α	14456	2	710	
14371	44739	Α	14457	349	493	SS*VRWVSGR*HPQ*STLSAPLG ETSGCHQGGLHDLKIVVL
14372	44740	В	14458	1	1338	
14373	44741	Α	14459	83	426	
14374	44742	Α	14460	93	345	APALCLQILLDCKSVRNRTSRF RPREVGIQQYKKGGGAAGKGG TGNAPGFLLFSKDS*KDPTPLPP PLLL*GQRDH*IQLHQE
14375	44743	Α	14461	1	1302	
14376	44744	A	14462	1	1174	
14377	44745	Α	14463	274 411	1428 627	HOLGURONITCHRONVINCORY
14378	44746	A	14464			HCLSHFQNTGPRGDVDFQQEL\ FPVLMADIRKEERSHLCRSSRR TWTILDKS*IQGSRLSSKEQGW GWGTSRK
14379	44747	В	14465	1	1553	
14380	44748	Α	14466	226	449	SLQPDI*CSGLRGSLLLV*PKSD
14381	44749	В	14467	I	2061	
14382	44750	В	14468	10	1099	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
14383	44751	Α	14469	34	406	GGGFISLDTGSHHKSHPGGS*H
						ALGWVRPHTLNSP\PSQQPGNC
		1	I	I		GNPAWEC*GPHG/APSRWSCST
						PCRPSHPKVPAVRVAGPAGGCT
						PYRT/PTRRTPR*RPGPCAACDR
						P\LHPHCGPRGPGAYPR
14384	44752	Α	14470	623	1671	VLRSVSSWLPTSMAMTVLAPM
1		1				LRAMSAGRLLM*PPSMSRCPSL
						GSHSGGR*PDS/GHAGAHVPPQ
		1	İ	l	1	GAIFVDSHG*GGDICGHVEVRQ
1		l	1		1	P/QGPQSAYPGRPDGCCG*RQT
		1		I		Q*TWR*LGT*GPTQVPAPKSCS
		1				WPTPGPAASCGWRCAPPRTTLR
1		l		١ ,	1	*KSQWTCRPPGPRGPPPPLMPL
		ŀ				VLRCASSPWPRPRLAQGQWCS
		l				P*APSPSGRSRCAGPRAWRKPS
1		1			ł	RRAHAAWQGCGAKRGGADSA
		1				GAGAPSTSCAYLATPRCAWGP
		l				HPPLGGAGLAA*GRAQPCSPAS
		1			ĺ	APAGRG*APGPPGPQ*GCSRPV
1				I		AGSARSRASARRPSGRVRYGV
1						HPPAGPATLTAGTLGCYPVRRP
						PGVPG
14385	44753	A	14471	32	972	QSGTFSTKNEGRTTRDVYDGR
		l		I		GPRVGRCGTCVYVRYVHDRPH
		l			1	APPHERARRNDGNTQRTGNAR
1		l				PGTAMPTATTDQPKGRSTRHK
		1		I		ETKRGRTARNGKQQNNPRK/RP
						RKDKRKQGSAGEGERGAGGKE
		1		I		KGQPRDTEAKGDRRTDFPPNG
		1				P*GQSKRGANIGGWFPPPRLGT
		1				PNPPLPGPQSAQVSGP*TPQMW
		1			1	MPTGTGCCGGSQ\SGSPPPWCV
						SP*PVAEALFHSST/PAPRHPSPP
		1				SSSPKHSHQPSTPPP*ISLGQGPC
		1				RPPKRRPAPDQPQVPRCPEPSC
				I		AYLATPRPTMRMQSACRGSAR
14386	44754	A	14472		3128	SRASARRPSG
14387	44755	В	14473	239	5474	
14388	44756	A	14474	542	781	PASIIMTGSISHITILTLRVNRLN
14,300	144,30	l^	14474	342	/61	ASIKRHRLTDWQIR/IKSQDLSV
		1				CYIQETHFTCKDPHKLKIKGWR
				I		KIYOVN*KOKKK
			Ь	L		hardry udayy

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14389	44757	A	14475	724	1340	QLTLQGRRVAPR*PSRVALEQE/ CGYNALLRYEGFENDSGLDFW CNICGSDIHPVGWCAASGKPLV
						PPRTIQHKYTNWKAFLVKRLTG
1				1		AKTLPPDFSQKVSESMQYPFKP
	İ			İ		CMRVEVVDKRHLCRTRVAVVE
				1		SVIGGRLRLVYEESEDRTDDFW
	ŀ			l		CHMHSPLIHHIGWSRSIGHRFK
		l		l		RSDITKKQDGHFDTPPHLFAKV
				l		RSFYKNPHL
14390	44758	A	14476	1822	2226	KRLFTNRTRITPFVAINSHHHHH
1.274						HHITUNTUTTTSTINSHHHHHY
				1		YHHHHHHHHHHOHTTTNTTT
		1				TNTTTTNNNTTTTITTTTTTTT
						TTSTINSHHHHHNQHCHHHHH
		į.			Į.	ннин/нининочнинин*нүо
						\ннниннүүннниннинно/
1				l		ннинионнимоннинооонн
ŀ		l				ДИННИАННОННИНОНАО /H
i i		l				ниннионсиннинининон
		l				ННОРРИННОН*НЅНОНН
14391	44759	A	14477	232	625	EGCCCIAIMVQKYQSPVRVYK
						YPFELIMALRK\RYPRCCVTCS*
		1				TGGAQQVGFRFLDLPLSLCTP*
		1				CLWRTSSSNF*RKAAGCSGSRL
1						QSHHFGLIPQPTAPKPWSSPLHI
		1				TKPQTLGQRDPGPRTRKRSK
14392	44760	Α	14478	1	808	MPKTTACNVHTVTWASCENAD
		l				SDSTGLRHHDSCEDPASPGRSD
						LCLVPHPTRPAVRGGRPAGGRC
		1				PAVPAEGPWRHLQQAQEGLKR
		l				LAGGLERLLENPVRRALPQGA
		l				QGHRAEPTEARPHPAPEHRRAG
		l				AAAERAERLESQPVGASAAAA
		l				SAASARIPYLVAVPAAGPPRHR
		l			+	CRRHRRGPGTSSGPRTALIRPAT
1		l				ARARAGIVSEPRTAPGATRARP
1		l			1	WRQGSHNPVAPSAP*LLRGPSIP
		1				WQIRPMPCPTSHQTCSERRPSG
1						RWQMPGSACRRSLETSSAGAG
l		1				RAQEAGGRLGETSGESG/RAGL
	1			I		YLKELRAIVLNQQRLVRTQRQS
				l		IDELERRLNELSA*NRSLWEHP
	1	1		i	1	QLPQAQPPPGFLTS*PSPLPALL
1	1		1	ŀ		GTAAAATAGAQEHLQDHGQRS
				l		SARPLPAPELASFQNHEQLLAQ
				l		PEPDPGGRAHTTQSPHQHPGTL
			1	l		GVKANKEKKGPPGCCAAPGTP
				l		LQQESPAPSARES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, *-possible nucleotide insertion)
14393	44761	A	14479	1	831	MIRYFGDYWTQALLTLIPGDPK
						HPRGPPVKVGVYGGQGYINSL
					ŀ	ALCHNFIWRELNRFSLLQDVPL
					l	VRYIDNIMLIGSIKFLGVQWCG
1						ACRDIPSKDPADPMVLEVSVAD
1						RDAVWSIWQALIDESQQRPLGF
		1				WSKSLPSSADNYSPFEROLLAY
		i				YWALVETERSTMGHQVTMLPE
1		1				LPVMNWVLSDPSSHK/ANGLA
ŀ		1			1	GWSGTGKKHDWKIGDKEIWRR
		1				GMWMDLSEWSK/D/VKIFVSHV
1				l		SAHQRVTSAEEEFNNQVDRMT
				l		RSMDTTQPLYPTTPVIAQWAHE
14394	44762	В	14480	56	3476	
14395	44763	Ā	14481	3	591	DPADPMVLEVSEADRDA\VPIS
1	111700	,		ľ.		ESOORPLGFWSKALPSSANNYS
	ļ					FFKROLLACYWVLVEIEHLTM
	İ					GHOVTMRPELPIINCVLSDPCSH
		1				KVGHAQQHSIIKWRWYIHDWA
1						EGTSKLHEEVAOIPMVSTPSLP
		1				OPAPMASWEVPYDOLTEEEKT
- 0				l		RAWFTDGSARHAGATOKWTA
						VALQPLSGTSLQDSSEEKSSQW
14396	44764	A	14482	1	1275	MEKNDIDQSRRRKVVEMNENV
1.1550	1	Ι΄.		ľ		QLVRYGEQASDLKTATKFKSPS
İ						EVKHIRFIQQAFIGSKLLMRPW
						TTVTKOILSLKSAKIALSESLIPC
						RKRAEFVEKQTQAFIMQVSDL
1	1					QQKVHAQPSQVSTVKVKALIG
1						KEWDPATWNGDVWEGPDEAG
						DTEFVNSDEAFLPEATAFPSPEV
					ŀ	GNDOTVWGLLDTGSELMLIPG
						DPKYYRGPPVKVGAYGGOVIN
						RGLAQIQLSMDPVGHRTRPVVS
1						SPLPECIIGTDILSSRONPHIGSLT
1					ł	DITLVHYIDDIMLIGSSEQEVTIT
1		İ				LDLLVQAVAQAALPVGPYDPA
	i					DPMELKVSVADRDAVWSLWQ
1						APIVTMRPERPIMNWVLSDPSS
						HKVGCAQQHSIIKWKWYIRDW
1						ARSGPEGTNHSRRVCS/LPQVQ
						APSAR*ATARRGQQAASGCCLF
1						WSCTCFSNFDV
14397	44765	A	14483	604	1760	NSWCRWFNFRY*YPTFYWES*
1,4397	177,00	ľ.		100.	1.,,,	NT*IPLREWFG*NHDASCP*TLD
1	1	1		I		SKSFSFETRTGSPCSSSLQTAYC
	1	1				GTLWIVQGV
14398	44766	A	14484	2710	3060	5.2, 40,
14399	44767	Ā	14485	3	227	
14400	44768	A	14486	3	266	
14401	44769	A	14487	2	564	
14401	1.7702		1. 7701	r <u> </u>	124.	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14402	44770	A	14488	2	503	FFFFLPIGFWRQHIPRLSVLLQHI Y*VT*NAASFEWGPEQEKAL*Q VQAVVQVALPLGPYDPADPML LEVSVADKDAGCTRL\QAPIGE SQWRSLGFRSKALRSSAENYSS FERQLLACYWALVETECLTLG HQVTI*PELPIMNWVLSDPSSHK VGHEOOHSIIKWK
14403	44771	A	14489	2	469	TOTIE QUIENTE
14404	44772	А	14490	Ī	633	MTVDYRKFNQVVTPMAA/AVP DAVSLLEQINTFPGTWYAAIDL NAFFSIPVHEAHOXQFAFULPQ GYINFPALCHNLIRRELDFFLLL QDITLVHYIDDILLIGSSEQEVV NTLDLLIHKSKEAEHTAASRIR VSCLPEQKSHEQTI.PWEQVPSS GDIKEYFPNAFVLLTTASLQGQ DNTSQLQLTWKAPEDIKMSKT DADADEEIEALRG
14405	44773	A	14491		518	MTVDYCKLNQVVIPIAAAVSD VVSLLEQINTSPGTWYAAIDLA NAFFSIPVHKAQQKQFAFSWQG QQYTFTVLPQWYINSPALCHNL IRRDLDCFSLPLDITLVHYIDDI MLIGSTIKWVVHSS/DSIIKWKW YVHDWARAGPEGTTNGLAG*S GTCKKHEWKTGDKGIRGRG
14406	44774	С	14492	33	932	
14407	44775	В	14493	27	2457	

1799

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14408	44776	A	14494	332	2158	RGYVFCSWKKTDGS/WRMTVD
		i				YCKLNOVVIPIAAAVSDVVSLL
						EQINTSPGTWYAAIDLANAFFSI
		1			1	PVHKAOQKOFAFSWOGOOYTF
						TVLPOWYINSPALCHNLIRRDL
						DCFSLPLDITLVHYIDDIMLIG\P
		l				ROLLACY/WALVETEHLTISHO
1		l				VTMRPELPIMNWVLFDPSSHKV
	1					GCAQQHSIIKWKWYVHDWAR
		l				AGPEGTTTPVISOWPHEOCGHG
		1				GRDGGYAWAOOCRLPLTKAD
						LNTATAKRPICQQQRPTLSPQY
		1				GTIPQGDQPATWWWVDYMGS
						LPSWKGQRFVLTGIDTYSGYGF
				1		AYPACNASAKTAICGLTECLIH
		1				HHDIPHSIASDOGTHFMAKEVR
						OWAHDHGIHWSYHVSHHPEA
						AGLIEWWNGLLKSOLOCOLGD
			İ			NTWQGWGKVLQKVVYALNQH
						PIYGTVSPIAKIHRSRNQGLEVA
		l				PLTITPRDPLAKFLLPFPATLOS
		1				AGLEVLVPEEGTLPPGDTMIPL
1		1				NWKLRLPPRHFGLLLPLNQQA
1				1	l .	KKGVTVLAGVTDLDYKDEITL
		1				LLHNGGKEEYAWNTGDPLGLL
		1				LILPCPMIKVNGKLQQLKPEAL
						VPKGVVFPPGDTTMLSLSWKL
	Ì	1				RLPSGHVGLLMPLSQQVQKGV
14409	44777	Α	14495	1	3187	MAEGKEEQVLSYTDGSRQREN
		1				EEDAKAETPDKTIRSHETYSLPR
		1				EWYEGNRPHDSITSQWVPPTTR
		l	İ			GNYGSTIQDEIWVGDHSGYVRP
				•		VPVPRSLNSDISYFGVGGKQAV
1						FFVGQSARMISKPADSQDVHEL
				1		VLSKEDFEKKEKNKEAIYSGYI
		ı				RNRKDDYDNHTGIDLVGTIIATI
		1		i		KGSNEEDTDTPLFIGKVRTLEFP
	1	1	l	1	l	FVNGSAEIMLMPSNQQHKTDE
		1		1		KGRANLGVFSVFAPRGEHTLQ
						VKAIYNKSIIEGP
14410	44778	Α	14496	266	467	
14411	44779	В	14497	152	1245	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino aeid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14412	44780	A	14498	1	1445	MLRAAAGGQGLAAAGMGLCA
14412	14700	ľ.		ľ		RPAGGSOLPRERMKGNRACLP
						LPPAVALTKGREKPDAGHQGR
						STOMMRESERGPCTHWTEERP
						GROLKILLOKOMRONTLMAFQ
					i	VSWDGRNKEIGRTFPHATAPHT
		1				QQLIFAMKIKRTGVTVEPWGLF
		1				ETGQRIRPIQAAGL*LSSSSTTC
		İ				VALAPLLTMHVKSLALLSAP/Q
1		1			l	MGALSTITSITTTRROTPAESHF
		1				PAARFHLLAPHSPTLPPCCPGEP
		1				SESAVTVLCFAKGKTAQDAAV\
		1				PTTSLTAAGHLAPRS*SSWSRM
						PVKVRRAPDSLPETSAHALPLT
						H*SWLPSASWGPASRALPGSEW
		1				VLKPVWQ/AITGPVSQITSSPGIP
ŀ						KP\GPLQA*GSAHPQALGFSVL
l						OIPLSNSPTSPMTPPLPRPKISNS
		1				SLPSQ/PPQALFPNRWTNSSSPV
i						HVITSAAQ\PRTPREWPAVSFYR
			1			TTATTASLIPHLWAPPQEN*QTS
		Ì				PAQNGSTLPCMSQATFHFPSP
14413	44781	A	14499	44	455	HLRNRTR/PSQITPHIYNHLIFDK
14413	4470:	,,	1,4427		1.00	/PLFNIWWWENWLAICRKLKLD
1		l				PFLTPYTKINSRWIKD*NIRPKSI
		l	l .			KNLEENLGNTIQHLG/TGKDFM
		1				TKTPKAMATKAKIDIWDLIKLK
	i i		1			SFCTAKETIIRVNRLPT*WEKIF
		1	1			AIYP
14414	44782	A	14500	1	346	
14415	44783	A	14501	1926	2448	DFIAVITTRKQLKTKYNAHHSK
1	1,,,,,,,	[`	1	1	-··-	OAITKHFSPKISDLILCLKKKIM
1		1	1	i		NRHFSKEDIYAAKKHMKKCSP
1		1		1		SLAIREMOIKTTMRYHLTPVRM
I			1			AIIKKSGNNRCRRGCGEIGTLLH
			1			CWWDCKLVQPLCKSMWRFLR
1		1		1		DL\DPAIPLLGIYPKDYKSCCYK
		1		1		DTCTRMFIAALFTIAKTWNQP
		_				D. C. C. C. C. C. C. C. C. C. C. C. C. C.

Jon., "possible nucleotide "ppossible nucleotide insertion) AEKAFDKIQQPFMLKTL GOTYLKHTRAIYDKPTA QKLEAFPLKTGTRQGGF NIVLEVLAQAIRQEKEI SKEEVKLSLFADDMILY. SAQKLLKLISLWSKVSKVSQ QKSQAFLYTNINQTESQ FTIATKRIKYLGIQLTEK EKYPELLNIKEDTINK
AEKAFDKIQQPFMLKTL JGTYLKITRAIYDKPTA QKLEAFPLKTGTRQGCI NIVLEVLAQAIRQEKEI JKEEVKLSLFADDMILY SAQKLLKLISNYSKVSG JKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRK
OGTYLKITRAIYDKPTA QKLEAFPLKTGTRQGCF NIVLEVLAQAIRQEKEI GKEEVKLSLFADDMILY SAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRD
OGTYLKITRAIYDKPTA QKLEAFPLKTGTRQGCI NIVLEVLAQAIRQEKEI GKEEVKLSLFADDMILY SAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESG FTIATKRIKYLGIQLTRE
QKLEAFPLKTGTRQGCF NIVLEVLAQAIRQEKEI GKEEVKLSLFADDMILY SAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRE
NIVLEVLAQAIRQEKEI GKEEVKLSLFADDMILY GAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRD
GKEEVKLSLFADDMILY SAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRD
SAQKLLKLISNVSKVSG QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRD
QKSQAFLYTNNRQTESQ FTIATKRIKYLGIQLTRD
FTIATKRIKYLGIQLTRE
KKYKPLLNKIKEDTNK
CSWIGRINIMKMAFPR
IENTWTQEGEHHTLGPV
KRGGIALVDIPNVNDKL
LARAIRQKKEIKGIQLG
LSLFADDMIVYLENSIV
KLISNFSKV\$GYKINVQ
LYTNNRQTESQIMSEFP
RIKYLGIQLTRDVKDLF
PLLKEIREDTNKWKNIP
INIMKMAILPKVIYRFN
PMTFFTELEKTTLKFIW
CIAKTILSKKNIAGGITL
YYKATVTKTAWYWYQ
OWNRTEASEVTSHIYNH
DKNKKWGNDSLFNKW
VLAICRKLKLDPFLTPYT
VIKDLNVRPKTIKTLEEN
DIGMGKDFMTKTPKA
KVDKWDVIKLKSFCTA
VSRQPTEWEKIFAIYPS
RIYKELKQIYRKK\TNN
AKNMNRHFSKEDIYAA
KCSSSLVIREMQIKTTM
FRFRKYREHHKDTPREE
*SSDSPKLK*RKKC*GQ
KLPTKGSPSD*KRISRQ/
RRQSWFFEKINKIDRPQ
KREKNQIDTIKNDKGD
EIQITIREYYKHLYANK
EMDKFLDTYTLPRLNQE
NRPITGSEIEAITNSLPTK
OGFTAEFYQRYKEELVP
OPIEKEGILPNSFYEASII
RDTTKKGNFRPISLMNI

SEQ ID NO:			SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	09/540,217	codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	,	١.		sequence	.,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	, , , , , , , , , , , , , , , , , , , ,
14418	44786	A	14504	1	2877	MENDFDELREEGFRQSNYSELR
						EDIQTKGKEVENFEKNLEECITR
	i					ITNTEKCLKELMELKTKAQELR
		1				EEWRSLRSRCDQLEERVSAME
		ı				DEMNEMKGEGKFREKRIKRNE
1		ı				QSLQEIWDYVKRPNLCLIGVPE
ŀ	1	l				RDGQNGTKLENTLQDVIQENFP
1	1	1				NLARQANVQIQEIQRTPQRYSS
1	1	1				RRATPRHIIVRFTKVEMKEKML
						RAAREKEIQTTISEYYKHLYTN
		1		1		KLENLEEMDKFLDTYTLPTLNQ
1		i		1		EEVESLNRPITGAE
14419	44787	Α	14505	I	2347	MELKTKARELHDECTSLSSRFD
		l				QLEERVSVMEDEMNEMNLPTK
	1	l				KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEPSII
1						LIAKPGRDTTKKENFRPISLMNI
		1				NAKILNKMLANQIQQHIKKLIH
	1					HDQVGFIPGMQGWFNIRKSINV
						IQHINRTKDKNHMIISIDAEKAF
		l				DKIQQHFMLKTLNKLVLEVLA
	İ				İ	RAIRQEKEIKGIQLGKEEVKVSL
		ĺ		İ	i	FADDMIVYLENPTVSAQNLLKL
	İ					IGNFSKVSGYKINVQKSQAFLY
1		l		i		TNNRQTERQIMSELPFTIASKRI
	1	l				KYLGIQLTRDVKDLFKENNKPL
1	1	l		l		LKEVKEDTNEWKNIPCSWVGRI
1		1				NIVKMAILPKVIYRFNAIPIKLP
		1				MTFFTELEKTTLKFIWNQKRAC
		1	İ	İ		IAKSIFSQKNKAGGITLPDFKLY
1				1		YKATVTKTAWYWYQNRDIAQ
		1				WNRTEPSEIMLHIYNYLIFDKPE
			İ	l		KNKQWGKDSLFNKWCWENWL
						AICRKVKLDPFLTPYTKMNSR
						WIKDLNVRPKTIKTLEENLGITI
				1		QDIGVGKDFMSKTPKAMATKA
	1			l		KIDKWDLIKLKSFCTAKETTIRV
				1		NRQPTTWEKIFATYSSDKGLISR
	1			1		IYNELKQIYKKKTNNPIKKWAK
	ļ					DVNRHFSKEDIYAAKKHMKKC
	1	ł				SSSLAIREMQIKTTMRYHLTPV
		1	l	l		RMAIIKKSGNNRKIQ/GGIWCD
			1	l		RIL*R*TTCRVAKEIQSL*RRI/W
14420	44788	В	14506	1	2265	KRLQRTLSIPVLDAV*PPMF*AS
14421	44789	A	14507	1	2832	
14421	44789	A	14508	1	2109	
14423	44791	A	14509	1	2757	
14424	44792	A	14510	i	2304	
14424	44172	<u> </u>	114310	ı <u>. </u>	2707	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
i				sequence		
14425	44793	Α	14511	li	2685	MGDFNTPLSTLDRSTROKVNK
	1	ŀ				DTQELNSAPHQADLIDIYRTLH
	1	l				PKSTEYTFFSAPHHTYSKTDHIL
		1				GSKALLSECKRTEHTNYLSDDS
						AIKLELRIKNLTQNRSTTWKLN
		1				NLLLDDYWVHNEMKAEIKMFF
	1					ETNENKDTTYQNLWDAFKAVC
		l				RGKFVALNAHKRKQGRSKIDT
		1				LTSQLKELEKQEQTHSKASRRQ
	1	l				EITKIRAELKEIETQKTVQKINES
	1	l				RSWFFERINKIDRQLARLIKKKR
		l				EKNLIDAIKNDKGDITTDPTEIQ
	1	l				TTIREYYKHLYANKLENLEEM
						DKFLDTYTLPRLNQEEVESLNR
		1				PITGSEIVAIINSLTTKKSPGPDG
		1				FTAEFYQRAIRQEKEIKGIQLGK
	1	1				EEVKLSLFADDMIVYLENPIVS
	i	1				AQKLISNFSKVSGYKINVQKSQ
		1				AFLYTNNRQTESQIMSELPFTIA
	1	1				SKRIKYLGIQLTRDVKDLFKEN
		ı				YKPLLKEIKEDTNKWKNIPCSW
	1	1				VGRINIVKMAILPKVIYRFNAIPI
		1				KLPMTFFTELKKTTLNFIWNQK
	1	1				\RAHIAKS/VLSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYQN
	1	l				RDTDQWNRTEPSEIMPRIYNYL
		1				IFDKPEKNKQWGKDSLFNKWC
	}	1				WKNWLAICRKLKLDPFLTPYT
	1	1				KINSRWIKDLNIRPKTIKTLEEN
		l				LGITIQDIGMGKDFMSKTPKAM
				l		ATKAKIDKWDLIKLKSFCTAKE
						TTNRVNRQPTKWEKIFATYSSD
14426	44794	В	14512	1	2337	

1804

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14427	44795	A	14513	1	1544	HINRAKDKNHMIISIDAEKAFD KIQQHFMLKTLNKLGIDGTYFR KSINVIQHINRAKDKNHMIISID AEKAFDKIQQHFMLKTLNKLGI DGTYFRKSINIDNPAYKQSQRQ KPLIISIDAEKAFDKIQQFFMLK TLNKLDNIIVYLENPIVSAPNLL KLISNFSKVSGYKINVQKSQAF KIKYLGIQLTRDVKDLFKENYK RIKYLGIQLTRDVKDLFKENYK RIKYLGIQLTRDVKDLFKENYK RIKYLGIQLTRDVKDLFKENYK RIKYLGIQLTRDVKDLFKENYK RIKYLGIQLTRDVKDLFKENYK
						PMTFETELEKTTLKFINVOKRA PVAKALSQKNKAGGITLPDFK LYYKATVTKTAWYWYONRDI GQWMRTEPSEITPHYYNYLIFD KPEKTKQWGKOSLFNKWEE NWLAICRKLKLDPFLTPYTKIN SRWIKDLMVRFKTIKTLEENLG TQDIGMGKDFMSKTFKAMAT KDKIDK WDLIKLKSFCTAKETT IRVNRQPTEWEKTFATYSSDKG LISRIYMEKLIYYKKTNYPIKK WAKDMNRHFSKEDIYAAKKH MKKCSSSLAIREMQIKTTVRYH LTPYRMAIIKKSGNN
14428	44796	В	14514	110	2153	DII VICIDINI CESSIII
14429	44797	A	14515		3210	MVKGSIQQEELTILNIYAPNTG ALRFIKOVLEDLORDLOSHTIIM GDFHTPLSTLDRSTRQKVNKDI QELNSALHQEOLIDIYRTLHPKS TEYTFFSAPHHTYSKIDHIVGSK ALLSKCKRTEIITNCLSDHSAIK LELRIKNLTONRSTTWKLNNLL LNDYWWHEMKA EIKMFFETN ENKDTTYQNLWDTFKAVCRGK FIALNAHKRUQERSKIDTLTSQL KELEKQETOPISKASRQEITKIR AELKEIETQ
14430	44798	A	14516	3	2820	ENKDITYONLIWDAFKAVCRG KFIALNAHKRKQERSKIDTLTS QLKELEKQEQTHSKASRRQEIT KIRAELKEIETOKSLOKINESRS WFFERINNIDRPLARLIKKKREK NQIDTIKNDKGDITTDPTEIQTTI REYYKHLYANKLKNLEEMDKF LOTYTLPRLNQEEVESLNRPITG SEIVTINSLPTKKSPGPDGFTAE FYGRYKEELVPFLLKLFQSIEKE GILPNSFYEASIILIPKPGRDTTK KENFR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14431	44799	A	14517	687	2274	LRDCKRINQHPVKTDQSAFCK
						M/API/MQDVVLEVLARAIRQEK
		1				EIKGIQLGKEEVKLSLFADDMI
l		l				VYLENPIVSAQNLLKLISNFSKV
		1				SGYKINAQKSQAFLYTNNRQTE
1		l				SQIMSELPFTIASKRIKYLGIQLT
	1	l				RDVKDLFKENYKPLLKEITEDT
		1				NKWKNIPCSWVGRINIVKMAIL
						PKVIYRFNAIPIKLPMTFFTELE
		1				KTTLKFIWNQKRARMAKSILSQ
	l	1				KNKAGGITLPDFKLYYKATVT
		ĺ	ļ			KTAWYWYQNRDVDQWNRTEP
		l				SEITPHIYNYLIFDKPDKNKQW
						GKDSLFNKWGWENWLAICRKL
						KLDPFLTPYTRINSRWIKDLNV
						RPKTIKTLEENLDITIQDIGMGK
l		l				DFMSKTPKAMATKAKIDKWDL
1		1				IKLKSFCIAKETTIRVNRQPTKW
		1				EKIFATYSPDKGLISRIYNELKQI
1		1				YKKKTNNPIKKWAKDMNRHFS
ł		l				KEDIYAAKKHMKRCSSSLAIRE
1		1				IQIKTTMRYHLTPVRMAIIKKSG
						NN/R/CF*WQKPGPSG*ALRSWK
}		1				EDIAGTSYVGMNQCQMRKNTK
ł		_				TLI
14432	44800	В	14518	1	3192	
14433	44801	Α	14519	1	2814	
14434	44802	A	14520	1	3144	MGDFNTPLSTLDRSSRQKVNK
ŀ		1				DTQELNSTLHHADLIDIYRTLHP
ł		1				KSTEYTFFSAPHHTYSKIDHVV
		1				GSKALLSKCKRTEIITNCLSDHS
1		l				AIKPELRIKKLTQNRSTTWKLN
		1				NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVS
						RGKFIALNAHKRKQKRCKIDTL
		1				ASQLKEVEKQEQTHSKASRRQ
				l	i	EITKIRAELKEIETQKTLQKINES
				1		RSWFLERINKIDRPLARLIKKKR
		\Box	L	<u> </u>	l	EKNQIDVIKNDK

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14435	44803	A	14521	1	2616	TTYONLWDAFKAVCRGKFIAL
						NVHKRKQERSKIDTLTSQLKEL
						EKQEQTHSKASRRQEITKIRAEL
						KEIETQKTLQKINESRSWFFERI
						NKIDRPLARLIKKKREKNQIDAI
						KNDKGDITTNPTEIQTTIREYYK
					-	HLYANKLENLEEMHKFLDTYT
					1	LPRLNQEEVESLNRPITGAEIVA
						IINSLPTKKSPGPDGFTGEFYQR
		l				YKEELVPFL/LKLFQSIEKEGI\M
						NIDAKILNKILANRIQQHIKKLI
						HHDQVGFIPGRQGWFNICKSIN
		1				VIQHINRAKDKNHMIISIDAEKA
		1				FDKIQQLFMLKTLNKLGIDGTY
						FKIIRAIYDKPTANIILNGKKLEA
						FPLKTGTRQGCPLSPLLFNIVLE
						VLARAIRQEKEIKGIQLGKEEV
1				1		KLSLFADDMIVYLENPIVSAQN
		ŀ		ļ		LLKLISNFSKVSGYKINVQKSQ
		l			1	AFLYTNNRQTESQIMSELPFTIA
						SKRIKYLGIQLTRDVKDLFKEN
						YKPLLKEIKEDANKWKNIPCSW
		ŀ			1	VGRINIVKMAILPKVIYRFNAIPI
					1	KLPMTFFTELEKTTLKFIWNQK
Ì		l		i	1	RAHITKAILSQKNKARGITLPDF
						KLYYKATVTKTAWYWYQNRD
l		l				IDQWNRTQPSEITPHIYNYLIFD
l		1	l	l	1	KPDKNKQWGKGSLFNKWCWE
l		1	1		1	NWLAICRKLKLDPFLTPYTKIN
						SRWIKDLNVRPKTTKTLEENLG
1		1	l	l		ITIQDIGMGMDFMSKTPKAMAT
ĺ		1				KDKIDKWDLIKLKSFCTAKETT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14436	44804	A	14522	782	3024	SKTVPTGNSLEPTQNCWLQQW LPQDIEIEKQEQT/HSKASRRQEI
						TKIGAELEEIETQKTLQKINESR SWFFEKINKSDRPLARLIKKKR
1						EKNQIDAIKNDKGDITTDPTEIQ
l		1				TTIREYCKHLYANKLENLEEMN
						KFLDTYTLPRLNQEEVESLNRPI
						TGSEIEAIINSLPTKKSPGPDGFT
	İ					AEFYORYKEELHINRTKOKNH
						MIISIDAEKAFDKIQQRFMLKTL
		l				NKLVLEVLARAIRQEKEIKGVO
1						LGKEEVKLSLFADDMIVYLENP
						IVSAONLLNLISNFSKVSGYKIN
	ŀ	l				VOKSOAFLYTNNROTESOIMSE
						LPFTIASKRIKYLGIQLTRDVKD
		1				LFKENYKPLLNERKEDTNKWK
	ĺ					NIPCSWVERINIVKMAILPKVIY
1	İ	1				RFNAIPIKLPMTFFTELEKTTLK
						FIWNQKRARIAKSILSQKNKAG
						GITLPDFKLYYKATVTKTAWYS
						YQNRDIDQWNRTEPSEILPRIYN
	ŀ			l		YLIFDKPEKNKQWGKDSLFNK
						WCWENWLAICRKLKLDPFLKP
						YTKIKSGWIKDLNVRPKTIKTL
						EENLGITIQDIGMGKDFMSKTP
1						KAMATKAKIDKWDLIKLKSFC
1				•		TAKETTIRVNRQPTEREKIFATY
						SSDKGLISRIYKELKQIYKKRTN
						NLIKKWVKDMNRHFSKEDVYA
				1		AKKHMKKCSSSLAIREMQIKTT
					i	MRYHLTPVRMAIIKKSGNNRITI
						VLLPGSLIVRSFHVTLNAILLHP
14437	44805	A	14523	1	1701	
14438	44806	Α	14524	1	4434	
14439	44807	Α	14525	I	2913	MGGPAALCIAVVPGDLVVALC
				i		LEEGLGSGIATWACSPWRISQG
	l					THLTFRSSLAVAAIPEGLPIVVM
						VTLVLGVLRMAKKRVIVKKLPI
			l			VETLGCCSVLCSDKTGTLTANE
1						MTVTQLVTSDGLRAEVSGVGY
	l					DGQGTVCLLPSKEVIKEFSNVS
1			1	l	1	VGKLVEAGCVANNAVIRKNAV
1						MGQPTEGALMALALKDQEDIY FMKGALEEVIRYCTMYNNGGIP
1			1	l		LPLTPQQRGAGPGDTALPSAAS
1						AFARMSAAERNDAFQG
L	L	<u> </u>	L	L	L	AFARINDAAERINDAFQG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14440	44808	A	14526	1	3009	MGDENTPLSTLDRSMRQKWKK DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL GSKALLSKCKRTEIITNYLSDGS AIKLELRIKNLTQNRSTTWKLN NLLLNDYWPPNEMKAEIKMFF ETNENKDTTYONLWDAFKAVC RGKFIALNAHKRKQERSKIDTL TSQUKELEKQEQTHSKASRRQE ITKIRAELKEIETQKTLQKINESR SWFFERNKIDRPLARLIKKKRE KNQIDTIKNDK
14441	44809	A	14527	1	4800	MGDENTPLSTLDRSTRQKVNK DTQELNSALHQADLIDIYRTLH DYSTEYTFFSAPHHTYSKIDHIL GSKALLSKCKRABIITNYLSDHS AIKLERIKNLTQSRSTTWKLN NLLLNDYWNNEMKABIKMFF ETKENKDTTYONLWDAFKAVC RGKFIALNAHKRKQERSKIDTL TSQLKELEKQEQTHSKASRRQE ITKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE KNQIDTIKNDK
14442	44810	В	14528	I	2901	`
14443	44811	A	14529	569	3443	IKH! TWKRTSSTSHHKNMPN- LEKQEQTHSKASRRQEITKIRAE LKEIETQKTVQKINESRSWFFER SNKIYRFLSRLIKKKREKNQIDT IKNDKGDTITDPTEIQTTIREFY KHLYANKLENLEEMDKFLDTY TIPKLNQEEIESLNRPITGTEIVT IINSLPTKKSPGPDGFTAEFVQR YKEELVPFLLKLFQSIEKEGILP NSFYEASILLAKPGRDTTKEK FRPISLMNIDAKILNKILAKRQ
14444	44812	A	14530	3591	8157	TGTSOKKTFMOPKKHMKKCST ITGHQSNANQNHNEIPSHTKLE WRSLKRSGNNRDVGNVVEAM YGGLPPPIMLIGHSMGGAIAVH TASSNLVPSLLGLCMIDVVEGT AMBALNSMONFLRGRPKTEKS LENAIEWRNIKDLSWADSIKNV AAASLACISGALPASAVTTSPD NKRMARNVLKVERLLAESPHH VVAEAVIORPNIPHLOTRDTYE GLCQTLGSQPTLYQIPSLYCSYE TNSNAYLLLOPIRKEVI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14445	44813	A	14531	172	328	AVKTEPEMKSOATKLSVVHEK
						KSOEGKPKEHTEPKSLPKOASD
						TGSNDAHNIKAVSR*SHROPSC
						LWFMRKNPKKESRKNTQSQKA
						YPSRHQIQEVTMLTI
14446	44814	Α	14532	3	660	TSPDCWEEGRSVDWPRPGPALS
						LHNCKLDLGIEGGTLQPPTPSF*
						PTGTRAV/PTRRRSRQETQLNSE
						RTVSPGSR**EKRTSRAW/RSPL
		1				QGPGPGHAGQVAGGTGPRTGS
						TRGD/TLQKCGKIQSVPLNCELI
		ı				PI*/PSILKPKVLPCSCPGIGASSS\
						CPSD*VFFCTS*PWPYLLSYPLS
l	+	1			1	MNFYRYHDHILKLLVQYTLYL
		ı				LLQFLQVLLKYLLSHPKQHS
14447	44815	Α	14533	1	1086	MVDNDVPWPLVVAQEKGQPD
						LCLSSKGYRPLALVQLDFHQVS
						VTSKIFILSGALRSLTGAATAD
					i	WTCRVLVIYPHDIMADSECNSL
						LVPSTTRKIMKHMLASEKIKTQ
1						NFNYNVDECVSLLYHHKANKL
1						EAAEARQECELLRVASVSRHC
1		l				W/RDPAHPHSCWPRC*APHCLG
ł						PARAPNARPAKPATTWNLRSSR
				1		APCAAPVPTQSWAGAPSGTAG
					l	AEPS*ECRPPENQARTAGSGVH
						AEACAPATCPACPGPGPCSGLP
						SSGRALFLL/CATLAIPCV*V*LC
						FLATPPPCR*HVTAPSAGKCIW*
						PDA*VTK/PRDLCGPVQVARDN
						GPRQVAQKKESSSYVSWFGTFI
						SRVGDRSPPPRRAATL
14448	44816	Α	14534	23	506	PDIDNIFSEFGAKSKPYLTVDQ
						MMDFINLKQRDPRLNEILYPPL
				1		KQEQVQVLIEKYEPNNSLARKG
1	1	l		1		QISVDGFMRYLSGEENGVVSPE
	1					KLDLNEDMSHPLSHYFINSSHN
l	1	1		1		TYLTGMNFLVLCYFSQSHQISP
	1	1				TSTFCCSEAEV/WFSSISCEVPLR
1	1	1		l		TLGFHH

SEQ ID			SEQ ID NO:			Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14449	44817	Α	14535	341	1248	KNAFTSFLTSFPPGQISVDGFMR
						YLSGEENGVVSPEKLDLNEDM
				1		SQPLCFMGIWGLFCWCFIAGQL
						AGNSSVEMYRQVLLSGCRCVE
				1		LDCWKGRTAEEEPVKILAY*IY
				1		MCSHF*EVIEAIAECAFKTLSLP
						TFLLLFLFSLSPKOQAKMAEYC
						RLIGSSIFKEYAFHFY*LESGVPL
						PSPMDLMYKILVKNKKKSHKS
			1			SEGSGKKKLSEOASNTYC*NGD
			l			MMICY*NLTGEADTESD\DADS
			1			GNV/CLFYSYFQGTAGSEAMAT
	1		ŀ			EEMSNLVNFLTFPFFVLLLLNSE
						RNKSFEMSSFVETKGL
14450	44818	A	14536	574	647	C*PGNM/HISGQHLVAGEKEAE
						SEDEEEEDMKLLGTPEGGSKFT
	i					OKKVKLAADEDDD/DDDADDD
			1			DDDDDDDDDDDDDEETEEKAPV
14451	44819	Α	14537	3	630	PGATHASAPPGVLRLKCGSGPV
						HIIGQHLVAVEEDAQSQDEEEE
						DVKLLSISGKRSAPGGGSKVPO
	1		1			KKVKLAADEDDDDDDDEEDDD
						EDDDDDDDFDDEEAEQQAPVKK
						SI\RDTPAKNAQKSNONGKDSK
						PSSTPRSKGQESFKKQEKTPKTP
						KGPSSVEDIKAKMQASI/EKKG
						GSLPKVEAKFINYVKN\CSRMT
			1		Į.	DQEAI\QDLWRWRRSL
14452	44820	Α	14538	1	1971	
14453	44821	Α	14539	316	1169	RVGGQSHGTQRISLCRHSVCSP
			1			LARQSPSNMK*VRTEEIQMAVS
			1			CYLKRWQYVDSDG\PLKQGLR
						LPQTAEELAANLTVHRESHCAD
						IVSAVP\CQAEPQQYEV\QFGRL
						RNFLIDSDSPHSHEVMPLLYPLF
						VCLHLSLVQNSPKSTVESFYSR
					1	FHGMFLQKAS\QKGVIEQLQTT
						QTIQDILSNFKLRAFLDNNYVV
						HLQEDSYNYLIHYLQSDSYTAL
						CKMFILLIHLDVQPAKTTD*QL
						YASGSSSRSENNYLEAPDMPSPI
						LONDAALEVLOERIKOVKDGC
	1	ĺ				PSLTTVCFYNT*QLLNTAEISPD
						SKLLAAGFDNSCIKLWSLQSRK
						LKSESCQVDMSRIHLACDILEE
14454	44822	Α	14540	1	297	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codnn for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14455	44823	Α	14541	2	1089	APLLGECRSGRGLASSRSRRRT
		ı				TMTQSVVVQGKLRRAGRDLVA
1		l				AESSGHVSCLMSYSLSFLSSGRS
1		l				RLALREHAAVNQVPARRREPPR
1		ı				LSARSSSPYSFHLQSE*L*KK1H*
1		1				LYFIFLSQAVLIDMEEGVVNEIL
1		1				L*IRTNQIKK*MQGIFKIHILIFR
		ı			İ	AVGHKVFGSLYQDQILEKFQIL
İ		ı				HNYILNHLC*FLKTL*SSGTGSG
		l		l		LGTFLLKVLEDEFPEVYRFVTSI
		ı	l	1		YPSGEDDVITSPFLVEYHLHPLF
		l		l		LIASLPMFFPQKSLFDIISKIDLM
		l				\VNSGKV/GGTTVKPKSL\VTSSS
1		l				GALKIELHLKCIILKLKIFSF*C*
1		l	l			LFSSARFEGSLNMDLNEISMNL
		1	l			VPFPPSS*EIVSLKYVLYFN
14456	44824	Α	14542	129	461	TCISGWHLFRASVTCPGIQTSPH
						NWHISDNSSLPNNIFIVCWNLL
		1				GMTRNYIIFIEQPLKMNLWKIA
		1				TSKIRGKAFSDGIS\GNPQCNTR
						FHVVEKRTGQLLPGRYYSKPF
14457	44825	Α	14543	1	522	
14458	44826	Α	14544	1	927	MEDLLTEARSPYWGRARAKIQ
		1				DSCWSVTSDPNAGSVDILAPLL
		1				LPSPSSYLPNFTCTVSVVTSHLH
		1				EAFLDLGFRGTGGSNCKTSRQV
]	l			QLNNQKRKKKKSKSLPFHMSL
		1				QIQNYKTQEEPHAKPSSQQHQH
		1				SEHESIQTKLVFRSSMTYMHTL
		1				KLIREKGSIMLSSTQGECPNSTC
						RWPAEEHPLGSTHREDQEHVIR
		ı				VPPEKVDLGETIHGVQVICSIAS
		l				TEKGKPSYYHSFGMTRNYIIFIE
		l				QPLKMNLWKIATSKIRGKAFSD
		l				GISWEPQCNTRFHVVEKRTGQT
		l				T*INPDKISIQKQHDIHAYIEAD
		I				KRKRKHNAVIHTG*M/YQI/GTC
		ı				RWPAEEHPLGSTHREDQEHVIR
	1	1			1	VPPEKVDLGETIHGVQVICSIAS
		1				TEKGKPSYYHSFGMTRNYIIFIE
		ĺ		1		QPLKMNLWKIATSKIRGKAFSD
		l			i	GISWEPQCNTRFHVVEKRTGQ\I
L		_		L	L	LNVKTMRMKTFMMIYYHLMN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14459	44827	Ā	14545	li .	882	MAEILTOVPDDMLKKKTPRVK
14439	44027	l^	14343	1	802	SVWRSQRGSCTPLIGTTELTLD
ŀ						TSHMSIRNMDRSHINVHTVRKP
		l				SADLPYFRTHEWAHTGGKPYD
						CEECGKSFISRSSIRRHRIMHSG
İ		l				DGPYKCNFCGKALMCLSLYLIH
						KRTHTGEKPYECKQCGKAFSH
		l	l			SGSLRIHERTHTGEKPYECSECG
		1	i			KAFHSSTCLHAHKITHTGEKPY
1		l				ECKQCGKAFVSFNSVRYHERT
	1					HTGEKPYECKQCGKAFRSASH
						LRTHGRTHTGEKPYECKQCGK
			1			AF*PRFQMTC*RRKLPE*NQCG
						EVSVGH\HSLNRHHRADTGHKP
		1				YEYQEYGQKPYKCTYCKKAFS
						YLPYFRTHEWAHTGGKPYDCE
		1				ECGKSFISRSSIRRHRIMHSGDG
1		l				PYKCNFCGKALMCLSLYLIHKR
1						THTGEKPYECKQCGKAFSHSGS
		1				LRIHERTHTGEKPYECSECGKA
		l				FHSSTCLHAHKITHTGEKPYEC
	İ	l				KQCGKAFVSFNSVRYHERTHT
						GEKPYECKQCGKAFRSASHLRT
		1				HGRTHTGEKPYECKQCGKAFG
						CASSVKIHERTHTGEKPCSSNTS
14460	44828	A	14546	3	395	SAEVGAAETTLTELRRTVQSLE
		ŀ				DLDSMRNLKASLENSL\GILLHL
	ľ					ESELAQTRAEGQRQAQEYEAL
						LNIKVKLEAEIATYRRLLEDGE
						DFNLGDALDSSNSMQTIQKTTT
						RRIVDGKVVSETNDTKVLRH
14461	44829	Α	14547	2	1044	AKTSSQMPSPQEGMCGKACTP
						ALSQADSLCPLLRLASEVEGYS
						LPACAEPYVQSECLSHLSVWSL
						QHALSSLASMSFTTCSAFTNYW
						SPGSVQVPSYGTQPVSHAASVY
						AGLGGSGSRISVSHFTNG\GGLA
		l				GMGGIQNEKETMQSLRDRLAS
						YLDRVRGLETENWKLESKIQEH
						LNVTRL*LETEIEALKKELLFM
1						KKNNEEEAKGLQAQIASSGLTV
l				1		EVDAPKS/QDLAKNMADSRAQ
1			1	1		YDKLARKNREELDKYWSQQIE
1		1		l		ESTRVVTTQSAKVGAAEMTLT
l		1				ELRHRVQSLEIDLDSTRNLKAS
1				l		LENSLREVEARYALQMEPSLLF
1				l		SKIIFGRVWEISYLHVFEGKQV
L		_		L	L	, some

SEQ ID SEQ ID NO: Met SEQ ID NO:			Amino acid sequence (X=Unknown,
	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	or pepade sequence	section, r possible national insertion,
1 1 1 1			
14462 44830 A 14548	1	1212	GPGTGIAGGLAGMGGIQNEKET
			MQSLNDRLASYLDRVRSLETE
1 1 1 1			NRRLESKIREHLEKKGPQVRD
			WSHYFKIIEDLRAQIFANTVDN
1 1 1 1			ARIVLQIDNARLAADDFRVKYE
			TELAMRQSVENDIHGLRKVIDD
			TNITRLQLETEIEALK*ELLFMK
			KTPEEEVKGLQAQ1ASSG\LTVE
			V\DAPKSQDLAKIMADIR\AQY
			DELARK\NREELDKYWSQQIEE
			STTVVTTQSAEVGAAET\TLTEL
			RRTVQVFGDSTWTSMRNLKA\
			NLENSL\GEVEARYAL\QMEQL\
1			NGILLHLESQLGQTPRTEAQRQ
			AQE\YEALL\NIKVKL\EAEICHL
			TRPPSWKIGEDFNL\GDS\LDER
		ļ	NSMQTIQKTTTRRISWIGQSGV
1 1 1 1			LRPIDTKVLEALSQQEASGTLL
			GKQEANKKFQSFK
	1	2382	
	2	173	
11100 11000 0 11001	1	732	AND ANALONO DA ANALONA
14466 44834 A 14552	455	682	AWKRAVALGKSRPMGAVAFK
			RRMQPVAGPRAMQEKPFPSSSS SLSIPPAF*YRLALSLAEPSRRW
1 1 1 1			RARTGALMOL
14467 44835 A 14553	158	368	KAKTGALMQL
	87	251	
	1	1026	
	201	578	
	336	767	AFVRAGSRFGGTPGARPDPAPA
		1	TRAPNPVGOSRHLRAASSRGRL
1 1 1 1			FLRRTRDRKFPRRPRLSRRLLG
		i	AI/GCRWNQPRAAEKSFSAGAV
			LDFSPGLSCLPAVQGSGPAARH
1 1 1 1			A*ASHPLHGLLCCPSIPDEHHPL
			LHGAQSHRPPKG
14472 44840 A 14558	6	535	ARHRVLIGVFTIPELDILSTSPPD
			SGAQLASPS\DPAPGLQVELPAS
			PARCARIPQPLGGRWDWAQTQ
1 1 1 1			EPSWLHLVEPAPGLQVELPASP
			TPCACIPQPLGDRWDWAPWSR
			GWCSLGRLGPHRSPWSGWEAE
			AWRAAGPEPCPTGRQLRLGEK
1 1 1 1			SSAAPVGWHCWGTQYTLRSH
1			WPGC
14473 44841 A 14559	375	638	
14474 44842 A 14560	92	514	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14475	44843	A	14561	54	380	SCHHYCRIHRRCARFLAGP*LPS RGARLGTCSPPCVSLPPAPWAP VRPEPPRRAPPPAPQRPVPSTTQ GLRSAGARRGTGRQLHLQLRG SAASFLKSVRPRTHQFRTH
14476	44844	Α	14562	53 I	773	
14477	44845	В	14563	92	1774	
14478	44846	Α	14564		390	LHDLQLSKPLSDDTALTDGG/Q PLPARWWAQPGRLQRCARFLA GP+LPSR/WGRARDLQPAMPEP PTPSVGSCAAQASPTSAAPCSM APSPIDHPRAWPVLCSFCMLLS LFVASISSCSQHQPMQVGTMVC L
14479	44847	A	14565	165	545	SLHSKLCWWLQQLWVTDTCPR CSVQDCRQRWASAGNQS*RLL VDPAPGLQVELPANPVPCARIP QPLGGRWD*APWSRGWCSSGR LGLHRSPWSGWEAQAWRAAG PEPRPAGROLGAMSCKVETGT
14480	44848	A	14566	1	309	SACLAGRPIRGEGRPRDLQPAM PEPPTPSMGSCAARASPTSATPC STAPSPIHHSRAEECGHTALG/H GRQLHLQPRNLPQTINWRTTYS SFPATLNRLDGGG
14481	44849	В	14567	1	843	
14482	44850	Α	14568	143	535	
14483	44851	Α	14569	375	571	
14484	44852	В	14570	289	972	
14485	44853	A	14571	27	363	
14486	44854	Α	14572	2	490	
14487	44855	A	14573	574	961	RLTLPDRLGSPPDTHQAQQITW ALLPQGFADSPHYFSQAQISSSS ITYLGIILHENTRALPADHV*LIS QTPISSTKQQLLSFLGMVRYFC LWIPSFTILTKPLY*FTKANLAD PTDPKSFPHSSFRSL
14488	44856	В	14574	182	1513	
14489	44857	A	14575	894	1305	GTOEPSWLHLVDPALGLQYELP ASPVPCARIPQPLGG*WDWAP WSRGWCSSGRLGPHRSPWSGW EAQA WRAAGPEPCPAGRQLRP GQHPL/RHMNKSQALSLLIAQP LGINLLSWWPPKYNRPVPLIGL SGLLIPATF
14490	44858	A_	14576	53 I	801	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14491	44859	A	14577	441	783	RAPPPAPWRPVPSTTOGLRSAG EQRGTGRQLHLSAAPVRDPRG EAHWAPESGRSAASILKPARPPA HREERTTPDAPP*EL*HSPRRSA ASLREDPWLHS*SQ*DQEPTNS GHTVSQDQSYYS*LEVSETKNP PIPDTL
14492	44860	A	14578	344	1632	FLAGPYLPSHGAGLGTCSLPCL SLPPTPWAPVREPPPMSTTPCS TAPSPIDHPRAEECERTAQDWQ AAPPAAPVWDPLGEASWAPES GTNISITPONATFYTEVQQQAW FASCITGHDMSSLKTSSVVVLR RQSEALLPANLTCNWQDSSAL ATLESALSOVRHKRFTFTLMVF TGSAIDIMATVSIAVVSITESVQ TTAFVDNLAKNICDELLLQDM DEAGNOHSQQNNTRTENQTPH VLTHKWELNNENIRIQGGEHYI LOPVGGRDENTLLRTYCLKSQ EGADTIKCRNKHNKETCKQSKS UYYSSTLNPEKLLVCKRLSLEL TRPENTLRCTNIGKLKAVAGG MSAASGIYLETDTQPYPLKQEL LSVSSTLNPEKLLVCKRLSLEL TREPENTLRCTNIGKLKAVAGG MSAASGIYLETDTQPYPLKQEL LSVSSTLNPEKLLVCKRLSLEL LSVSCYAFARCLDAKHSTSENV ELGKEKCVGKNSVVVVFLVRK LSVSQVAFARCLDAKHSTSENV ELGKEKCVGKNSVVVVFLVRK LSVSQVAFARCLDAKHSTSENV ELGKEKCVGKNSVVVVFLVRK LSVSCYAFARCLDAKHSTSENV

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop cndon, /=possible nucleotide deletion, \=possible nucleotide insertion)
1	sequence		05/340,217	sequence	or peptide sequence	deterior, i-possible macreonae miser non)
]				oequeue-	1	
14493	44861	Α	14579	3	2233	NWLPSVSRCQETHKIAANQSLT
						QRQMPLTSLYCDVVILPVSVGL
						ESLOSPYYRTSPCDPVSQFCLIK
						ISASSLPVLSLWKTLKNAQSYQ
1						RDYRHDNKAHYFQKLVTMQE
						NLYVLLCDEELDAVSLEAVHQ
						QKMKNRQAHSSSGCCQKVAN
						APRNEESSGQFRVIGNEKFEKG
	İ					PCGVRQQIFEKVLLPNRTCEFV
						TPRFNRNFEDLIKDLKAAETLID
						FLVYEPKNQRRLVNRFRLPCKL
						LHNLGHPLKTDAAQSPRKPPGP
		l				SWMPSFGSKPPSHPPLVSPHLNP
						QVWDTSSPSLATEHASLTISLKP
						NHPYPAQSQYSIPQHALKGSKP
1						VITRLLEHGLLKPINSPYNSPILP
		1		l		VQKLDKSYRLVQHLCLIYQIVL
1	į.				1	PIHPMVPNPYTLLSSIPFTTHYS
			ļ			VLDLKHAFFTIPLYPSSQPLFAF
						TWTDPDTHQAQQITLAVLPQCF
1			1			IDSPHYFSQAQISSSSVTYLGIIL
		l				MKTHIGLGAVEQGVVLVGEAR
					1	AAQEPMEWVGGSGM/CGLQVP
						SPAPWEGS*GPARNRAQ\PVTIV
						PVLDFNPAFHIIPDTTPDHHDCI
	}	ļ		i		SLIHLTFTPFPHISFFPVPHLEHT
		ŀ				WFIDGSSTRPNCHSPAKAGYAI
	1	l				VSSTSIIEATALPRSTTSQQAKLI
		1				AYTDSKCAFHILHHHAVIWEER
					İ	NFLTMQGASIINAFLIKTLLKDI
						LLPKEAGVIHCKGHQKASDPIT
						QDNAYADKVAKKQLAFQLLSL
14494	44862	Α	14580	1	642	SGTVPPTFVRTVIQAQGARTVI
						QQSLIGSLYIFRHCVRCQGIESV
						FVIIIWSRVGYDEKGFCLFTSVL
		İ				GVWGHVRTKPCLLITWRTLKS
						RSGLE*PAGFSSSPGTSPQIIRKQ
		ĺ				KRGVPQHPVCDLPGRFRPRAGT
	1					PLAAQIQVPGLPPCGHQE*NRR
1			1			WAGSSALCPPYLCARAGTGAV
					1	CRACEPAGV/AGWAWAWWAP
		L				HSEHA ASPAGPRQWGT
14495	44863	В	14581	393	549	
14496	44864	В	14582	I	591	
14497	44865	С	14583	226	828	
14498	44866	Α	14584	1	1362	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \range possible nucleotide insertion)
14499	44867	A	14585	1	585	MANVHQTMPMRLEKLPDSFKP PEPKSNSQRVSTEAGTAGALTP QHVRAHSSPASLQLGAVSPGTL TPTGVVSGPAATPTAQHPQQSS FEIPDDVPLPAAREMAKTSSGQ RYFLNHIDQKTTQQDPRKTML SQMNVIANPPVQQNMMNSIPQS PQGGV/MGGSNSNQQRQMRPQ QLQMEERLQLKQQ*LLQQELA LR
14500	44868	Α	14586	345	1053	RWRLQNLSFCSSCSGGSGAT CFARGRQSNLQSKIKFPSNTD SRSTGYNSFDRQPNICFYKRQE TPHTGPIIPVPEGRPYMPG*CG A*SQFPASRN*SEKN*SYCH HTAFNVIESSVI**RYCTRQWHI VKCLDSFCDPFLISDELRRASTD AGTAGALTPGIVRAHSSPASLQ LGAVSPGTLTPTGVVSGPAATP TAQHLRQSSFEIPDDVPLPTSW GEAKTSGGSKIFP
14501	44869	В	14587	607	1185	

1818

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino aeid sequence (X-Unknown,
NO:	of peptide		in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
ŀ			İ	sequenee		
14502	44870	Α	14588	1	2142	MIILIDAEKAFDKIOOPFMLKTL
14302	44070	^	14366	ľ	2142	NKLGIDGTYLKITRAIYDKPTA
						NIILNGOKLEAFPLKTGTROGCP
						LSPLLFNIVLEVLAQAIRQEKEI
						KGIOLGKEEVKLSLFADDMILY
						LENPIVSAOKLLKLISNVSKVSG
		i I				
			l			YKINVQKSQAFLYTNNRQTESQ
						IMSEFPFTIATKRIKYLGIQLTRD
			1			VKDLFKKYKPLLNKIKEDTNK
			l			WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
ŀ			1			VGWGKRGGIALVDIPNVNDKL
		1				MVLEVLARAIRQKKEIKGIQLG
						KEEVKLSLFADDMIVYLENSIV
1			l			SAQNLKLISNFSKVSGYKINVQ
1			ŀ			KSQAFLYTNNRQTESQIMSEFP
1			l		8	FTIATKRIKYLGIQLTRDVKDLF
1						KENYKPLLKEIREDTNKWKNIP
1						CSRIGRINIMKMAILPKVIYRFN
						DIPIKLPMTFFTELEKTTLKFIW
1			l			NQKRACIAKTILSKKNIAGGITL
						PDFKLYYKATVTKTAWYWYQ
1			ł			NRDIDQWNRTEASEVTSHIYNH
			1			LIFYKPDKNKKWGNDSLFNKW
						CWENWLAICRKLKLDPFLTPYT
			ł			KIHSRWIKDLNVRPKTIKTLEEN
						LGNTIQDIGMGKDFMTKTPKA
						MATKAKVDKWDVIKLKSFCTA
			l			KETTIRVSRQPTEWEKIFAIYPS
						DKGLISRIYKELKQIYRKK\TNN
						PIKKWAKNMNRHFSKEDIYAA
						NRQMKKCSSSLVIREMQIKTTM
14503	44871	В	14589	1	1647	
14504	44872	В	14590	1	2025	
14505	44873	В	14591	4	2121	
14506	44874	A	14592	2	1879	
14507	44875	A_	14593	1	2130	
14508	44876	Α	I4594	I	3018	
14509	44877	В	14595	1	3171	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14510	44878		14596	l .	2700	MEKKOSRKTGNSKNOSASPPP
14510	448/8	A	14396	I	2700	
						KECSSLQATEHSWMENDFDEL
		l				REQGFRRSNYSELKEEVRTHGK
						EVKNLEKRLDEWLTRITNAEKS
						LKDLMELKTMARELLDECINLS
						SQFDQLEERVSVMENQMNEMK
İ						PNLRLIGVPESDGENGTKLENT
						LQDIIQENFPNLARQANIQIQEIQ
				i		RMSQRYSSRRAIPRHIIVRFTKV
						EMKEKMLRAAREKGRVTLKG
						KPIRLKADLSVETLQARREWGP
		ŀ				IFNILKEKNFQPRTSYPGKLSFIS
		ŀ				EGEMKSFTDKQKLRYFVTTRP
						ALKELLKEALNMERNNRYQPL
i						EKQIQTTIREYYKHLYANELEN
				ļ		LEEMDKFFDTYTLPRLNQEEVE
						SLNRPITGFEIEAIINSLP/TKKSP
1						EPGGFTAEFYQ\RIEKEGILPNSF
						YEASIILMLKPGRDTTKKENFRP
		l				ISLMNIDAKILNKILANRIOOHI
		l	ŀ			KKLIHODOVGFIPGMOGWFNIR
		l				KSINIIOHINRTNDKNHKIISIDA
		l		ŀ		EKAFYKIQQPFMLKTLNKLGID
		l				GSYLKIIRAIYDKPTANIILNGQ
						KLEAFPLKTGTROGCPLSPLLF
		l				NLVLEVLARAIRQEKEIKGIQLG
		1				KEEVKLSLFADDMIVYLENPIA
						SAONLLKLKGNFSKVSGYKINV
						OKSOAFLYTKNRQTESOIMSEL
ı						PFTIASKRIKYLGIQLTRDMKDL
						FKENYKPLLNEIKEDTNKWKNI
	l				ŀ	PCSWEGRISILQMAILPEVIYRF
14511	44879	_	14597	1	2310	T CS W ESIGNIEQ WATER EVITIG
14511	44880	A	14598	1	1305	
		Λ				
14513	44881	A	14599	1	2148	
14514	44882	В	14600	1	777	
14515	44883	Α	14601	2	2971	
14516	44884	A	14602	1	3810	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14517	44885	A	14603		1773	MNIGAKIINKILANQIQHIRKLI HHDQVGFIPGMQGWFNICKSIN VQHINKTKOKNHMIISIDAEKA CDKIQQRFMLKTLNKLGIIDGKY LKIIRANYDKIPANIILINGQKLE AFPLKTGTRQGCPLSPLLFNIVL EVLARAIRQEKEIKDIQLGKEEK KISLCADKMIVYLENPIVSAQN LLKLISMFSKVSGYKINVQKSQ AFLYTNINRQTESQIMSELPFTILA KSRIKYQJQHQTRDVKDLFKEN YKPLLKEIKEDTINKINIPCSCVG RINIVKMAILPKELEKTTLKFIW NQKRARITKSILSQKNKAGSIM LPDFNLYYKATVSKTAWYWY QURDIDQWWRTEPSEITPHIYN YLIFDKPEKNKQWGKDSLFNK WCWENWLAICRELKLDPLTL YTKINSRWIKDLNVSPKTIKTLE ENLDITIQDIGMGKDFMSKTAK MAMAKAMSTKAKFGKWDLL LKSFCTAKETTIRVNIRSILK STSLQAFSRKILNCSKRFFKLAQ MCKLVELRPQMHITSRGSGNC SEARTLLCDFSAGRTSVNHGRI WROIREMGADPEIGLEVCNINN
						PTSFTLFYFLSYTTG
14518	44886	В	14604	1	2070	
14519	44887	В	14605	I	3190	
14520	44888	Α	14606	1	3370	

1821

14521 44889 A 14607 569 3030 RKH*TWKRTSSTSHHKNMFN	SEQID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
14521	NO:		hod				
14521		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
LEKQEGTHISKPSRREITKIRG ELKEIETPKTAQKINESRSWFS RINKIDRPLARLRKKKEKKOR DTIKNDEGDITTHPTEIHTIRE YKHLYANKLANLEEMOKFLD YTLPRPNQEEVESLINGPIAGSI QAIINSLPTKKSPGPPDGFYQRY EELVPFRLKI-FQSIEKEGILPN YEASILLIPKORDTTKKENPR SILLINIAKILIKALIANIQQHII KLMHHDQVGFIPGMQGWFNI KSINVIQHINRTKDKNHMIISII AERAFDKIQPFMLKTLINKLL DGTYHKIIRANYDKPTANILIN QKLEAFPLKTGTQGCLPSEIL FNIVLEVLARAINGEKETEGIQ GKEEVKLSLFADDMIVYLEN VSAQNLLKLISNFSKVSGYKI VQKSQVFLYTDNRQTESQIISE PFFTIASRIKKYLAQDITYVXKU FKERYYMEIKEDTNKWKNIPCS WVGRINIVKMALIPKVIYRRN SIQLPMTFFTELEKTTLKFIWN KRAHIAKSILSKNKAGGIML DFKLYYKATVTKTAWHVYY RDJDQRNRTEPSEIMPHVYNH FGKFDKNRQWGNDSLFNKW WENNLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 I 2742 14524 44891 A 14609 I 3828 14524 44891 A 14610 I 2667					sequence		
LEKQEGTHISKPSRREITKIRG ELKEIETPKTAQKINESRSWFS RINKIDRPLARLRKKKEKKOR DTIKNDEGDITTHPTEIHTIRE YKHLYANKLANLEEMOKFLD YTLPRPNQEEVESLINGPIAGSI QAIINSLPTKKSPGPPDGFYQRY EELVPFRLKI-FQSIEKEGILPN YEASILLIPKORDTTKKENPR SILLINIAKILIKALIANIQQHII KLMHHDQVGFIPGMQGWFNI KSINVIQHINRTKDKNHMIISII AERAFDKIQPFMLKTLINKLL DGTYHKIIRANYDKPTANILIN QKLEAFPLKTGTQGCLPSEIL FNIVLEVLARAINGEKETEGIQ GKEEVKLSLFADDMIVYLEN VSAQNLLKLISNFSKVSGYKI VQKSQVFLYTDNRQTESQIISE PFFTIASRIKKYLAQDITYVXKU FKERYYMEIKEDTNKWKNIPCS WVGRINIVKMALIPKVIYRRN SIQLPMTFFTELEKTTLKFIWN KRAHIAKSILSKNKAGGIML DFKLYYKATVTKTAWHVYY RDJDQRNRTEPSEIMPHVYNH FGKFDKNRQWGNDSLFNKW WENNLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 I 2742 14524 44891 A 14609 I 3828 14524 44891 A 14610 I 2667	14521	144880	Δ	14607	569	3030	RKH*TWKRTSSTSHHKNMPN*
ELKEIETPKTAQKINESRSWFS RINKIDRPLARLRKKREKNQ DTIKNDEGDITTHPTEIHTIIBE YKHLYANKLENLEEMDKFLD YTLPPROGEVESLINGPIAGSI QAIINSLPTKKSPGPDGFYQRY EELVPFRLKI-FQSIEKEGILPNS YEASILLPRYGROTTKKENSRI SLLNINAKILNKILANRIQQHI KLMHDQVGIPFAGQWFM) KSINVIQHINRTKDKNHMIISII AEKAFDKIQQFPMLKTLNKLC DGTYHKIIRANYDKPTANILLI QKLEAFPLKTGTRQGCPLSPLI FNIVLEVLARAIRQKEETEGIQ GKEEVKLSLFADDMIVYLENN VQKQVFLYTDIRQTESQIISI PFTTASKRIKYLGIQLTRYVKI FKERYNBIKEDTNKWKNIPCS WVGRINIVKMAILPKYLGRYN. SIQLPMTFFTELEKTTLKFIVN KRAHASSLISKNAGGIML DFKLYYAATVTKTAWHVYQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNRQWGDSLFNKW WENVLAICKLKLDPFLTPYI KINSRWIKDLNYRPETIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 114524 44890 A 14609 1 3828 114524 44890 A 14610 1 2667 114525 44893 A 14611 1 2325	14321	44007	n	14007	1507	3030	
RINKIDEPLARIRKKKEEMOQ DTIKNDEGOITTHPTEIHTITIEE YKHLYANKLENLEEMOKFID YTLPRPNQEEVESLINGPIAGSI QAJINSLPTKKSROPDGFYORY ELVPFRLKI-POSIEKEGII PNI YEASIILIPKPROTITKKENFRI SLLNINAKILKNILANRIQQHIII KLMHHDQVGFIPGMQGWFNI KSINVIQHINRITKOKNIMIISIII AEKA-FPLKITGKOKNIMISIII QKLEAFPLKTGTRQGCPLSPLI FNIVLEVLARAIRQEKETEGIQ GKEEVKLSLFADDMIVYLENB VSAQNLLKLISNFSKVSGYKIN VQKSQVFLYTDNRQTESQIISE PHTIASKRIKYLGIQLTRYVKL FERRYNEIKEDTNKWKNIPCS WVGRINIVKMALIPKVIYRRN. SIQLPMTFFTELEKTTLKFIVN KRAHIAKSILSKKNKAGGIML DFKLYYKATVTKTAWHWYQ RDIDQRNRTEPSEIMPHVYNHI FGKPKNNQWGDSLFNKW WENVLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LONNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14609 I 3828 14524 44891 A 14609 I 3828 14524 44892 A 14610 I 2667							
DTIKNDEGOITTHPTEHTIIRE YKHLYANKLENLEEMDKFLD YTLPRPNQEEVESLINGPIAGSI QAIINSLPTKKSPQPDGFYQRY EELVPFRLKI-PSSIEKEGIL.PN YEASIILIPKIGGDTTKKENPRI SLLNINAKILIKKLANRIQQHI KLMHHDQVGFIPGMQGWFNI KSIMVQHINKTKOKNHMIISII AEKAFDKIQOPMIKTINKLC DGTYHKIIRANYDKPTANILIN QKLEAFPLKIGPMIKTINKLC DGTYHKIIRANYDKPTANILIN QKLEAFPLKIGPTRQCCPLSPII FNIVLEVLARAIRQEKETEGIQ GKEEVKLSLFADDMIVYLENE VSAQNILKLINSFKSVGSYKI VQKSQVFLYTDNRQTESQIISE PFITIASKIKYLQGLTRYVKI FKERYNEIKEDTNKWKNIPCS WVGRINIVKMAILPKVIYRFN, SIQLPMTFTELEKTTLKFIVM KRAHIAKSILSKKNKAGGIML DFKLYYKATVTAWHVYQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNQWGNDSLFNKW WENWLAICKLKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMRKDFTSKTFRA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14610 1 2667 14525 44893 A 14611 1 2325		1					
VKHLYANKLENLEEMDKFLD		į .	l				
YTLPRPNQEEVESLINGPIAGSE QAIINSLPTKKSPGPDGFYQRY EELVPFRLKI-FQSIEKEGIL-PN YEASILIPKPGRDTTKKENPR SILLNINAKILIKALIKANIQQIII KLMHHDQVGFIPGMQGWFNI KSINVIQHINRTKDKNHMIISIE AEKAFDKIQQPFMLKTLNKLC DGTYHKIIRANYDKPTANILIN QKLEAFPLKTGTQGCLPSEIL FNIVLEVLARAINGEKETEGIQ GKEEVKLSLFADDMIVYLEN VYAQNILKLISNFSKVSGYKIN VQKSQVFLYTDNRQTESQIISE PFFTIASKIKYLGQLTRYVKL FKERYYBEIKEDTNKWKNIPCS WVGRINIVKMAILPKVIYRFN. SIQLPMTFFTELEKTTLKFIWN KRAHIAKSILSKNKAGGIML DFKLYYKATVTKTAWHVYQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNRQWGNDSLFNKW WENNLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14609 1 3828 14524 44892 A 14611 1 2325		1					
QAINSLPTKKSPGPDGFVQRV		1					
EELVPFR.ki.pgsiekegil.nn YEASIILIPKPGROTTKKENFRI SLLNINAKILANRIQQHII KLMHHDQVGFIPGMQGWFNI KSINVIQHINRTKDKNHMISIII AEKAFDKIQQPFMLKTLNKLC DGTYHKIRANYDKPTANILN QKLEAFPLKTGTRQGCPLSPLI FNIVLEVLARAIRQEKETEGIQ GKEEVKLSLFADDMIVYLEN VSAQNLLKLISNFSKVSGYKIN VQKSVFLYTDNRQTESQIISE PFITIASKRIKYLGIQLTRYVKL FERYYBEIKEDTNKWKNIPCS WVGRINIVKMALIPKVIYRN. SQLPMTFFTELEKTTLKFIVN KRAHIAKSILSKKNKAGGIML DFKLYYKATVTKTAWHWYQ RDIDQRNRTEPSEIMPHVYNHI FGKPDKNRQWGDSLFNKW WENWLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LONNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKKLKSFCT 14522 44890 A 14608 1 2742 14510 14610 1 26567 14523 44892 A 14610 1 2325 145216 145215 14893 A 14611 1 2325 145216 145216 145216 1 2325 145216 1 2325 145216 1 2325 1 145116 1 2 2325 1 145116 1 2 2325 1 145116 1 2 2325 1 145116 1 2 2 2 2 2 2 2 2 2		1					
YEASILIPKPGRDTTKKENPRI SLLNINAKILNKILANRIQQHB KLMHHDQVGFIPGMQGWFNI KSINVIQHINRTKDKNHMIISII AEKAPENLQOPFILKTINKLC DGTYHKIIRANYDKPTANIILN QKLEAFPLKTGTRQGCPLSPLI FNIVLEVLARARIQEKETEIGI GKEEVKLSLFADDMIVYLENP VSAQNLLKLISNFSKVSGYKIN VQKSQVFLYTDNRQTESQISIE PFITASKRIKYLGIQLTRYVKI FKERYMEIKEDTNKWNNIDES WVGRINIVKMAILPKVIYRFN, SIQLPMTFFTELEKTINKENING KRAHIAKSILSKKNKAGGIML DFKLYYKATVTKTAWHVQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNKQWGNDSLFNKW WENWLAICKKLKLDPFLTPYI KINSRWIKDLNYPETIKLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14524 44890 A 14610 1 2667 14525 44893 A 14611 1 2325 14525 44893 A 14611 1 2325		1	l				
SLLNINAKILNKILANRIQQHIK KLMHHQVGFIPGMQGWFNI KLMHHQVGFIPGMQGWFNI KLMHHQVGFIPGMQGWFNI KSINVQHINKTKOKNHMISISI AEKAFPKIQOFPMLKTLNKLC DGTYHKIRANVAPEPTAINLE GKEFVLSLFADDMIVYLEN VOKSQVFLYTDNRQTESQISE PFITASKRIKYLGIQLTRYVKI FERRYNBEIKDTNKWKNIPCS WVGRINIVKMAILPKVIYRRN. SIQLPMTFFTELEKTTLKFIWN KRAHAKSLISKKNKAGGIML DFKLYYKATVTKTA WHWYQ RDIDGRNTEPSEIMPHVYNH FOKPOKNRQWGDSLFNKW WENWLAICKLKLDPFLTPYI KINSRWIKDLNYPEPTITLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14609 1 3828 14524 44892 A 14610 1 2667 14525 44893 A 14611 1 2325 SILNINAKILKKILANRIQQFIP KENNHQLINAKILANRIQCE KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KENNHQLINAKILANRIQQFIP KINSRWINGLINAKILANRIQQFIP KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILANRIQQE KENNHQLINAKILAN		1					
KLMHHDQVGFIRGMQGWFNI KSINVIQHINRTKDKNHMIISIE AEKAFDKIQQFFMLKTLNKLC DGTYHKIIRANYDKPTANILIN QKLEAFPLKTGRQGCLPSIL FINVLEVLARAIRQFKETEGIQ GKEEVKLSLFADDMIVYLENF V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLLKLISNFSKYSGYKIN V9AQNLIKLISNFSKYSGYKIN V9AQNLIKLISNFSKINGLISNFSKYSGYKIN V9AQNLIKLISNFSKYSGYTANI V9AQNLIK V9AQNLIKLISNFSKYSGYKIN V9AQNLIKLISNFSKYSGYKIN V9AQNLIKLISNFSKYSGYKIN V9		ł					
KSINVIQHINRTKOKNIMIISII		1	1				
AEKA-PÓKIQOPPMLKTLNKLC							
DGTYHKIIRANYDKPTANILIN		1	l				
QKLEAFPLKTGTRQGCPLSPLI			l				
FINIVLEVLARAIRQĒKETĒGIQ GKEEVKLSLFADDMIVYLENF VSAQNLLKLISNFSKVSGYKIN VORSQVELYTDNRQTESQIISE PPFTIASKRIKVJGQLTRYVKL FKERYNEIKĒDTNKWKNIPCS WVGRINIVKMAILPEVIYRRN, SIQLPMTFFTELEKTTLKFIWN KRAHIAKSILSKNKAGGIML DFKLYYKATVTKTAWHWYQ RDIDQRNRTĒPSEIMPHVYNH FGKPDKNKQWGNDSLFNKW WENNLAICKKLKLDPFLTPYI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMKGPTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14512			l				
GKEEVKLSLFADDMIVYLENE VSAQNLLKLISNFSKVSGYKIN VQKSQVFLYTDNRQTESQIISE PFTIASKRIKYLGIQLTRYVKL FEERYMEIKEDTNKWKNIPSE PFTIASKRIKYLGIQLTRYVKL FEERYMEIKEDTNKWKNIPSE WVGRINIVKMAILPKVIYRFN. SIQLPMTFFTELEKTTLKFIWN KRAHIAKSILSKKNKAGGIML DFKLYYKATVTKTAWHWYQ RDIDQRNRTEPSEIMPHVYNH FOKPDKNKQWGNDSLFNKW WENWLAICKKLKLDPFLTPYI KINSRWIKGLNYPETTIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT LGNNIQDIGMGKDFTSKTPKA LGNNIQDIGMGKDF			1				
VSAQNLLKLISNFSKVSGYKIN			l				
VQK\$QVFLYTDNRQTESQIISE			l				
PFTIASKRIKYLGIQLTRYVKI FKERYYNEIKEDTNKWKNIPCS WVGRINIVKMALIPKVIYRRN. SIQLPMTFFTELEKTTLKFIVN SIQLPMTFFTELEKTTLKFIVN KRAHAKSLISKKNKAGGIML DFKLYYKATVTKTAWHVYQ RDIQRINTEPSEIMPHVYNH FCKPDKNRQWGDSLFNKW WENVLAICKLILDFTLTYT KINSRWIKDLNYBPKTIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14514 14614 14615 14							
FKERYNEIKEDTNÄWKNIPCS							
WYGRNIVKMAILPKYIYRFN, SIQLPMTFFTELEKTILKFIWN KRAHIAKSILSKKNKAGGIML DFKLYYKATVTKTAWHWYQ RDIDQRNRTEPSEIMPHYYNH FOKPDKNKQWONDSLFNKW WENWLAICKKLKLDPFLTPYI KINSRWIKDLNYPKTIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890							,
KRAHIAKSILSKKNKAGGIML DFKLYYKATVIKTA WHWYQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNKQWGNDSLFNKW WENWLAICKLKLDFFLTPYI KINSRWIKDLNVRPKTIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDK WDLMKLKSFCT			l				WVGRINIVKMAILPKVIYRFNAI
KRAHIAKSILSKKNKAGGIML DFKLYYKATVIKTA WHWYQ RDIDQRNRTEPSEIMPHVYNH FGKPDKNKQWGNDSLFNKW WENWLAICKLKLDFFLTPYI KINSRWIKDLNVRPKTIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDK WDLMKLKSFCT			1				SIOLPMTFFTELEKTTLKFIWNO
RDIDQRNRTEPSEIMPHVYNH FGKPDKNRQWGNDSLFNKW WERWILAICKKLKLDPFLTPTI KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14610 1 3828 14524 44892 A 14610 1 2667 14525 44893 A 14611 1 2325			l				KRAHIAKSILSKKNKAGGIMLP
FGKPDKNKQWGNDSLFNKW WENWLAICKKLKLDFFLTPYT KINSRWIKDLNYPETIKTLE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14610 1 3828 14524 44892 A 14610 1 2667 14525 44893 A 14611 1 2325							DFKLYYKATVTKTAWHWYON
WENWLAICKILKLDPETITYT KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT MATCDKIDKWDLMKLKSFCT MATCDKIDKWDLMKT MATCDKIDKWDLMKT MAT			l				RDIDORNRTEPSEIMPHVYNHLI
KINSRWIKDLNVRPKTIKTLEE LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890			l				FGKPDKNKOWGNDSLFNKWC
LGNNIQDIGMGKDFTSKTPKA MATKDKIDKWDLMKLKSFCT 14522 44890 A 14608 1 2742 14523 44891 A 14609 1 3828 14524 44892 A 14610 1 2667 14525 44893 A 14611 1 2325							WENWLAICKKLKLDPFLTPYT
MATKDKIDKWDLMKLKSFCT			l				KINSRWIKDLNVRPKTIKTLEEN
14522 44890 A 14608 1 2742 14523 44891 A 14609 I 3828 14524 44892 A 14610 I 2667 14525 44893 A 14611 I 2325			l				LGNNIQDIGMGKDFTSKTPKA
14523 44891 A 14609 I 3828 14324 44892 A 14610 I 2667 14525 44893 A 14611 I 2325			l				MATKDKIDKWDLMKLKSFCTA
14524 44892 A 14610 1 2667 14525 44893 A 14611 1 2325	14522	44890	Α	14608	1	2742	
14525 44893 A 14611 1 2325	14523	44891	Α	14609	I	3828	
	14524	44892	Α	14610	1	2667	
14526 44894 A 14612 1	14525	44893	Α	14611	1		
	14526	44894	Α	14612	1	3057	

ASAADTQANRVWSGPPANSN PAAIRVLTVRRKTNKQKGHPH QNPICTSPSSKTERTQTTIREYY KHLYANRLENLEEMDKFLDTT TLPRLNQEEVESLNRIITESDH AINSLPTKSSOPPOFTAEFYY RYKEELVPFLLKLPQSIEKEGH PNSYFASHLIPLKI GROTTKEK NPRPISLMNIDAKIL NKILANRI QQHIKKHIHQQVGFIPGMQGW FNIRKSINVIQHINKTNOKNIM VSIDAEKAFDKIQQPFMLKTLN KLGHOFPYLKIRATYDKLTAN LNGQKLEAFPLKTGTRQGCLL PLLFNIVLEVLARAIRQEKEIK QLGKEEVKLCLFADDMIVYLE NPIVSAQNLLKLISNFSKVSGY INVQKSQAFLYTNRQTESQH SELPFTIASKRIKYLGQLTGSON SELPFTIASKRIKYLGQLTGSON SELPFTIASKRIKYLGQLTGSON KDFYKENYKPLLNEIKEDTDK: KNINCSNVGRINMIKMALIPKY YRRNAIPIKLPMTFTFLELKTTI KFIWNQKRARIAKSILSQKNK GGITLPPKLYYKATVTKTAC RVGRAQQHSISKWKWYHDW QVGPGTNIDSARYPDTTOKW AAALQPLSRTSLKDSHEGKSS WAELRAVHLVLRFAWEKWE	SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino neid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
WAELRAVHLVLRFAWKEKWI DVQLYTDSWAVASGLAGWSC TWKKHDWKIGDKEIWGRATP IAQWAHEQRGHGGRDGDYAW 14528 44896 B 14614 1 3105	14527	44895	A	14613		2973	DNTARGTIRQOHLLFTNIRCSA ASAADTQANRVWSGPPANSNR PAARVLTVRRKTNKQKGHPH QNPICTSPSSKI KERTQTTIREYY RHLYANRLENLEEMDKFLDTY TLPRI.NQEEVESLNRRITESDIE AINSLPTKSSPGDGFTAEFYQ RYKEELVPFLLKLFQSIEKEGIL PNSFYEASIILIPKLGROTTKKE PSIRKSINVIDAKILINKILANRI QOHIKKHIHQQVGFIPGMQGW PSIRKSINVIQHINRITNDKNHMI VSIDABEKAFDKIQOFPMLKTLN KLGIDGPYLKIIRAIYDKLTANI LNGQKLEAFPLKTGTRQGELLS PLLFNIVLEVLARAIRQEKEIKGI QLGKEEVKLCFADDMIVYCLFADDMIVYCLFADMIVYCLFADMIVSU SELPFTIASKRIKYLGIQLIGDV KOPFRENYKPLLNEIKEDTDKW KNIPCSRVGRIMIKMAILDKVI YRFNAIPKLPMTFFTELEKTIL KFIWNQKKARIAKSILSQKNKA GGITLPPFKLYYKATVTTACH RVGRAQQHISIKWKYHIDWS GOGTIPPFKLYYKATVTTACH RVGRAQQHISISKWKWHIDWS KOYGPGGTNOSARYPDTTOG W
14528 44896 B 14614 1 3105							WAELRAVHLVLRFAWKEKWP DVQLYTDSWAVASGLAGWSG TWKKHDWKIGDKEIWGRATPV
	11500	11005	_	14614	ļ.——	2106	IAQWAHEQRGHGGRDGDYAW
			-				

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ĺ	sequence		
14530	44898	Α	14616	735	2839	RSFPRSRPSPFLLLSRYLRIHMV
						FSVLPFGLQNPKYLLSGSLQEK
			ĺ			FRTPGINSHKTLDPNRVIIKVRR
	l	l	•			RLEKEKALRAYVGKSEVRNMH
			ŀ			LRERCKL*KKREKNQIDTIKND
		l				KGDITTDATEIQTTIREYYKHLY
						ANKLENLEEMDKFLDTYILPRL
			l			NQEEVESLNRPITGSEIEAIINSL
						PTKRSPGPDGFTAEFYQTYKEE
		l				LVPFPLKLFQSTEKEGILPNSFY
						EASIILIPKPGRDTTKKENFRPIS
						LMNIDAKILNKILANRIQQHIKK
						LIHYDQVGFIPGMQGWFNKRK
	ŀ					SINVIQHINRTNDKNHMIISIDAE
						KAFDKIQQPFMLKTLNKLGIDG
			l			TYLKVIRAIYDKPTANIILNGOK
	l		ł			LEAFPLKTDTROGCPLSPLLFNI
						VLEVLARAVROEKEIEGIOLGK
						EEVKLSLFADDMIVYLENPIVS
						AQNLLKLISNFSKVSGYKINIQK
İ						SQAFLYTNNRQTESQIMSELPFT
						TASKKIKYLGIQLTRDVKDLFK
						ENYKPLLNEIKEDTNKWKNIPC
		1				SWVGRINIMKMAILPKVICRFN
						AIPIKLPMPFFTELETTTLKFIWN
1			ł			
			l			QKRARIAKSILSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYQN
						RDIDQWNRTEPSEITPHIYNYLI
		ŀ				FDKPEKMSIIDTGGWYEAPVSS
		l				FFKEGLCRTGTENQVNRQPPAA
						SPFKVYVSCTGSLHPRPHYFLE
14531	44899	A	14617	1	2250	
14532	44900	A	14618	1	2870	MKAEIKVFFETNENKDTTYQNL
1						WDTFKAVCRGKFIALNAHKRK
			l			QERSKIDTLTSQLKELEKQEQT
			ŀ			HSKASRRQEITKIRAELKEIQTQ
1						KTLQKINESRSWFFERINKIDRS
						LARLIKKKREKNQIDTIKNDKG
		1	l			DITTOPTEIQTTIREYYKHLYAN
1			l			KLENLEEMDKFLDTYTLPRLNQ
			l		ĺ	EEVESLNRPITGAEIVAIINSLPT
			l			KKSPGPDGFTAEFYQSWAETQP
1		l	1			KKENFRPISLMNIDAKILNKILA
		l				KRIOOHIK
			L		L	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino neid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14533	44901	A	14619	248	2623	RQWAGVVGRCSHLASWVSSN
	1				ł	SETGAIRSSTEVDAPDDSMLLS
		ŀ	1			CDIDLTAARRAWLGCLPTKKS
			l			GPDGFTAEFYQRCKEELVPFL!
						KLFQSI/EKEGILPNLFDEASIIL
		Į.				PKRGRDTTKKENFRPISLMNIE
						AKILNKILANRNQQHIKKLIHF
		ļ				DQVGFIPGMQGWFNICKSINV
	1					QHINRTKDKNHMIISIDAEKAI
						DKIQQPFMLKTLNKLGIDGTY
						KIIRAIYDKPTANIILNGQKLEA
						PLKTGTRQGCPLSPLLFNIVLE
		1	1			LARAIRQEKEIKGIQLGKEEVK
						SLFADDMIVYLENPIVSAONLI
						KLISNFSKVSAYKINVQKSQAI
						LYTNNRQTESQIMSVFPFTIAS
	i					RIKYLGIQLARNAKDLFKENY
	1					PLLNEIKEDTKKWKNIPCSWV
	1					RINIVKMAILPKVIYRFNAIPIK
	1				ŀ	PMTFFTELEKTTLKFIWNOKR
	1					RIAKSILSQKNKAGGITLPDFK
						HYKITVTKTAWYWYQNRDID
						WNRTEPSEITPHIYNYLIFDKPI
						KNKQWGKDSLFNKWCWENW
						AICRKLKLDPFLTPYTKIHPRW
					1	KDLNVRPKTIKTLEENLGNTIC
	1			1		DIGMGKDFMSKTPKAMAAKA
	1	ı				KIDKWDLIOLKSFCTAKETTIR
						NROPTKWEKIFATYSSDKGLIS
	1					RIYKELKOIYKKK\TNNPIKKW
						AKDMNRHFSKEDIYAANKHM
						KKCSPSLAIREMQIKTTMRYH
14534	44902	A	14620	1	3253	MGDFNTPLSTLDRSTROKVNK
14334	44902	^	14020	1	3233	DTQELNSALHQGDLIDIYRTLI
	1		1			PKSTEYTFFSAPHHTYSKIDHI
						GSKALLSKWKRTEIITNYLSDI
				1		SAIKLELRIKNLTQSRSTTWKL
						NLLLNDYWVHNEMKAEIKMF
	1				i	
			1			ETNENKDTTYQNLWDAFKAV
	1.	1			l	RGKFIALNAHKRKQERSKIDTI
		1	1			TSQLKELEKQEQTHSKASRRQ
			1			ITKIRAELKEIETQKTLQKINES
						SWFFERINKIDRPLARLIKKKR
		L_		ļ		KNQIDTIKNDK
14535	44903	В	14621	I	2758	
14536	44904	A	14622	I	5178	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	Incation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14537	44905	A	14623	1	3382	MGDFNTPLSTLDRSTRQKVNK
		-		1		DTQEFNSALYQADLIDIYRTLH
		ŀ				PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCERTEIITNYLSDHS
		l				AMKLELRIKNLTQNCSTTWKL
		l				NNLLLNDYWVHNEMKAEIKM
						FFETNENKDTTNONLWDAFKA
1						VCRGKFIALNAHKRKQERSKID
ŀ						TLTSOLKELEKOEOTHSKASRR
1						QEITKIRAELKEIETQKTLQKIN
						ESRSWFFERINKIDRPLARLIKK
						KREKNQIDTIKNDK
14538	44906	Α	14624	965	4091	TWKGTTSTSRCKIMPKYRSTRQ
						KVNKDTQELNSALHQADLIDIY
		1				RTLHTKSTEY/TFFSAPHHTYSK
		l				IDHIVGSKALLSKCKRTEIITNY
i		1				LSDHSAIKLELRIKNLNQSRSTT
		1				WKLNNLLLNDYWVHNEMKAE
l				l		IKMFFETNENKDTTYQNLWDA
ŀ						FKAVCRGKFIALNAHKRKQERS
		1	1			KIDTLTSQLKELEKQEQTHSKA
1		1				SRRQEITKIRAELKEIETQKTLQ
		1		1		KINESRSWFFERINKIDRPLARLI
						KKKREENQID
14539	44907	A	14625	3	3229	
14540	44908	В	14626	1	2791	NAME OF THE PROPERTY OF THE PAR
14541	44909	A	14627	1	3158	MVKGSIQQEELTILNIYAPNTG
						APRFIKQLLSDLQRDLDSHTLIM GDFNTPLSTLDRSTROKVNKDT
		l				QELNSALHQADLIDIYRTLHPK
	1					STEYTLFSAPHHTYSKIDHILGS
						KALLSKCKRTEIITNYLSDHSAI
ļ.						KLELRIKNLTOSRSTTWKLNNL
i						LLNDYWRKQERSKTDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETOKTLOKINESRSWFF
1						ERINKIDRPLARLIKKKEKNOID
						TIKNDKGD
14542	44910	В	14628	I	7849	
14543	44911	Α	14629	1	3985	MENDFDELREEGFRRSNYSEL
1		1				WEDIQTKGKEVENFEKNLEECI
		1				TRITNTEKCLKELMELKTKARE
		1				LHEECRSLRSRCDQLEERVSAM
						EDEMNEMKGEGKFREKRIKRN
1						EQSLQEIWDYVKRPNLHLIGVP
	1					ESDGENGTKLENTLQDIIQENFP
1	1	1				NLARQANVQIQEIQRMPQRYSS
						RRATPRHIIVRFTKVEMKEKILK
	1	1	1			AAREKDRSTRQKVNKDTQELN
		1	1			SALHQADLIDIYRTLHPKSTEYT
						FFSAPHHTYSKID
1						FFSAPHHTYSKID

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
14544	44912	В	14630	64	5735	
14545	44913	В	14631	891	6894	
14546	44914	A	14632	33	476	
14547	44915	A	14633	1	360	GOGEPKOLGRHPGVLRPHVVS
1				ľ		AKTGDAC/IWTOVRNLFPERPK
						VQWKEGCA\LQSGNSQESVTEQ
						GSKVSAYSLSSPLTLSKADYEK
						HKLYACEVTHOGLSSPVTKSFN
						RGEHEGEVPPCI
14548	44916	A	14634	40	508	LPGECHRAGQQGQRLQPQLLG
,	,					LLMLWVPGSGGEIVMTOTPLSS
1						SVTLGQPASISCRSSQSLVHSDG
						NTYLTWLOORPGOPPRPLIYHV
						FKRFSGVPDRFSGSGAGTDFTL
						E/ISGRVEA*\DVGVYYCMQG\T
						OFPCSFGQGTKLETKRTVAAPS
14549	44917	Α	14635	3	652	AWWNSETPAQLLFLLLLWLPY
						TSGEIVLTQAPGTLSLSPGERAT
						LSCRASOTIGSTYLAWYOORPG
						QAPRFLIYGASSRATGIPDSSSSS
				<u> </u>		SSSSSSSSSSSSSSSSSSSQQ
1	ŀ	l				YNNWPPGGTFGP\GTKVEIKRT
						VGCTHLFFIFPAHLMSQLKIWE
						LPSACVPVCNNF\YPREVQSYSG
1						KVDN/VPSNRVTPRESVTEQD\S
				ł		KDSTLQPSAATLDA
14550	44918	Α	14636	3	440	
14551	44919	Α	14637	457	1270	SRRVSFPPSA*QAVSDPSRQQLP
			}		İ	SLRLRWGPEGRSFPAENLGRGV
						RWVGGRDPLAA\LP*SSGPLNA
						GRLFARPQESGAPRIGNVRDVP
						RAREDVGTQAGRREREFPRLFG
			i			PAAVVGCERHLSPFPRAAGAG
	!					GRAGAAGRRSAPPGGLGSCRSC
						RISAGPHGNDVPTLGGRGW/PL
						*NSGAEKL*EQ*G*/TGTGSWSL
						AAILGGVPREP\GFRGAGRETLG
	İ	l				PRGWLVMLSRAVLLHSVLLCT
	l	l				CHSSWVLLHSDLACHADNNPD
				L		IQSCLFPMSKY
14552	44920	С	14638	90	465	
14553	44921	Α	14639	116	439	
14554	44922	С	14640	278	364	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14555	44923	Α	14641	116	687	EIKDAALQLPLWRAQDRERAM
l						KERREKRREREKERDREKKRK
						EKMFCKRSCHTIMEPTPELSCV
						LPCLAPIVLAPPPLPL\PVLPLPY
		i				ATR*LPQQNRQSTLTEAHGL\TR
						GPITLATNRCMPSTTRDRRRLT
		1				PPSTLPNTTNLSWGVREGPNQD
						PEPSLSPCVPHDPFDYNAESLFS
						YWLVSVTPMTAHTP
14556	44924	Α	14642	474	1166	HPNRQGLRAPATAPGHPGREQ
						ALCEAPQGPGQKCGPPRAQEA
						Q/GTEPRRASRAAGACRHHWP
		ļ.				QVA\RPGAGAPAGAARLGGV*E
		1				PGRAA/APRLPLPSQCCHR*GTS
		l				PAHLSASRSSSATCPWRRTAGR
						SGPSPCMTL\KQRQGHPRDGWI
		l				RAPSCDVTNRGLSSLLNLGGSA
		ı				QQLLLASRPLGSGSGALEALEH
		l				GKSRAPWEPWLHGTHNPVHGK
		<u>_</u>				DPAGSTSHWPRPPAPRT
14557	44925	A	14643	283	806	PRGSFSSTGRTRNAAGSTSNCSL
1		l				RSLSARIEERKVSRVCGVPNHR
		l				HLRLGALTTTRPSWNGAIPTLA
		l				KLPAKRC*RRSGRRSKTWSFPG
		1				ST*LGNFAEEMICAASARPAGV
		l				PGPAL\PTPLRAPLPGVPWTSSR
		1				GRMCARLSCC/HAGRPVPARPG
14558	44926	A	14644	1	338	AARARRAAAGERHRPPALRV
14559	44927	Ā	14645	1	1091	MSNDGRSRNRDRRYDEVPSDL
14337		ľ`	1.10.15	ľ	1.07.	PYODTTIRTHPTLHDSERAVSA
		1				DPLPPPPLPLQPPFGPDFYSSDT
		1				EEPAIAPDLKPVRRFVPDSWKN
		l				FFRGKKKDPEWDKPVSDIRYIS
İ		i i				DGVECSPPASPARPNHRSPLNS
		l				CKDPYGGSEGTFSSRKEADAVF
		l				PRDPYGSLDRHTQTVRTYSEKV
		l				EEYNLRYSYMKSWAGLLRILG
		l				VVELLLGAGVFACVTAYIHKDS
			l	1		EWYNLFGYSQPYGMGGVGGL
1	1		İ	1		GSMYGGYYYTGPKTPFVLVVA
		1				GLAWITTIIILVLGMSMYYRTIL
			1	i		LDSNWWPLTEFGINVALFILYM
					i '	AAAISLM*MIPTRGGLCYYPLF
			l	l		NTPVNA\GSAG*KEDR*LQ*SSC
		1		l		LSP**FISLVLWFA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nuclentide deletion, *=possible nucleotide insertion)
14560	44928	A	14646	398	1107	SPELVCGLLIFCKYTYPICAGPY RCPFGGLRH*SAR*QPPSGLREN GARPFGGEWLCPC*FPP*PLNSR WKLPPRLVT*GEPWAAGEKFP RLGACTGPAAGGLPRARDWE PGRAGPERWAPGROPAGTPAK TAAVKRSAPLPGSGODYANGER KPPSPPCQWSGNL*VLCG*RL SLPAA*KPKKA*RRGRQSPGK TGYQVQOTRPKNDPKSSIGPSP KNRQSLCLQPLWSFLIQ
14561	44929	Α	14647	1	540	MGVRDPLEEAVCALAELQHCA GRFAALFRASRGERISLIKLCP QPLPPRALSQGYKPLIGILLAF FQRCTAQRGGI*RGSLATVTLIS CGGLCPV*TSQRRCCLPVRGKLP TQASVMANAPPRTRLQHLRSTS DCCAGSENFKPVDLSLLGSVGV GSTELDHLAPWLQPAFQGSEW FYLTF
14562	44930	Α	14648	169	487	EVCYYPPSETYFCQFVKLILCPV LFPCWQGVVIQPKLSCMRCLST \LLRGVSQ*GYMGVRDPLEEAV CPLAELEHCAGRSAALFRAGRQ ERLSLLKLRPQLPLSPSA
14563	44931	Α	14649	1	210	
14564	44932	В	14650	1	1683	
14565	44933	A	14651	1	529	MDSEVQADDEVSDENGELIGN WGKGGMGVSDPLEEAVCPLAE LERSAGRSAALFRASRQECLSL LKLT/LTAAPSPRGTAFRSPFKS PAQRRSIIC*Q/LYSCKTFRSTVA PMPRLCLPQASGKVQ*EN*SWA IPA*QGTSPMCSLCSGAPHQPG APLVLPLPPPVNRLRRELQCWL
14566	44934	С	14652	295	430	
14567	44935	A	14653	452	637	
14568	44936	A	14654	151	446	MVLSHWHSRCHWGMKKNYLQ LARCLPKWLPSFVFETQGPCDV STQGTLLVYGLQRPWEKHSI*A R*HHPSQHGPSQLPLARGGSSP TPCASRMRQHPT
14569	44937	A	14655	2	335	EDRSAFRPRQPHTLHPLHARSL APRSPTPPSPPSPDTQLGLSGPTS GPESAPTA/PGNPSWRSSRWGSS SPCAASST*KSPYP*/CSPT/CAFP SPRLPFCRSAYQPAAGAGRGK
14570	44938	В	14656	373	502	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14571	44939	A	14657	1	400	GSPAPAAMDEERALYIVRAGE AGAIERVLRDYSDKVKSPOPGI AVSHPRAALRACTSHPSPPSF GCALVPPKTFPAWACVFVPPA SIHAS/SHPTTPTVAPTPLIPGPA A*PPVLLSSVQVTPPESVDVHL
14572	44940	A	14658	659	1177	ARKGSALLRLYLSNAWCVCVI VLRIFSTASAYITRRDSSLGPV TFLNFSFGSPGEIPTPFFFPVFP LPLVAWFCPASFCLPQCPLLVP: EPSPPPHSGDAHCSPPKTFPAW ACVFVPPAFSIHAS/SHPTTPTV ACVFVPPAFSIHAS/SHPTTPTV APTPLIPGPAA*PPVLLSSVQVT PPESVDVHLCPVPHS
14573	44941	В	14659	1	318	
14574	44942	Α	14660	2	131	
14575	44943	А	14661	354	762	CPEIRPPS*THGTK/SHFPPTASC THSVLEAL/PDTQRQPSPRVHQ DPSPAUPALLPSPHEHGPQHLLL QLPSHRCHPDTAPFNCDAVKA PGHATAALEHPYLSGRQSPPSF THLPKLPAGRGYTVCALPKGP ERSAV
14576	44944	А	14662	157	413	ALVCS/SSLAIREMQIKTTMRY LTPVRMAIIKKSGNNRCWRGC GEIGTLLHCWLDCKLVQPLWK SVW*FLRNLELEIPFDPAIPLL
14577	44945	A	14663	1926	2448	DFIAVITTEKQLE TKYNAHHSK QAITKHFSPKISDLILCLKKKIM NRHFSKEDIYAAKHIMKKCSP SLAIREMQIKTTMRYHLTPVRN AIIKKSGNNRCRRGGEIGTLLI CWWDCKLVQPLCKSMWRFLR DL\DPAIPLLGIYPKDYKSCCYK DTCTRMFIAALFTIAKTWNQP
14578	44946	A	14664	1	1584	
14579	44947	Α	14665	1	4729	
14580	44948	Α	14666	1	1566	
14581	44949	A_	14667	1	1424	
14582	44950	A	14668	1	576	
14583	44951	A	14669	1	3325	
14584	44952	A	14670	1	2195	
14585	44953	A	14671	1	3162	
14586	44954	В	14672	287	2018	
14587	44955	IA	14673	11	1073	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
14588	44956	Α	14674	1	3335	MVKGSIQQEELTILNIYAPNTG
						APRFIKOVLSDLORDLDSHTLI
						MGDFNTPLSTLDRSTROKVNK
						DTQELNSALHQADLIDIYRTLH
1	İ					PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
1		1				AIKLELRIKNLTQSRSTTWKLN
		1				NLLLNDYWVHNEMKAEIKMFF
		1				ETNENKDTTYQNLWDAFKAVC
						RGKFIALNVYKRKQERSKIDTL
	l					TSQLKELEKQEQTHSKASRRQE
		l				ITKIRAELKEIETQ
14589	44957	A	14675	1	1838	MIIPIDAEKAFDKIQQPFMLKTL
	1					NKLGIHGMYLKIIRAIYDKPTA
1	1					NIILNGQKLEAFPLKTGTRQGCP
1	İ					LPPLLFNIVLEILARAIRQEKEIK
		1				GIQLGKEEVKLSLFADDMIIYLE
		1				NPIVSAQNLLKLISNFSKVSGYK
		1				INVQKSQAFLYTNNRQTESQILS
	1	ł				ELPFTIASKRIKYLGIQLTRDVK
						DLFKENYKPLLNEIKEDTNKW
						KNIPCLSIGKINIMKMAILPKVIY
	ł			İ		RFNAIPIKLPMTFFTELEKTTLK
						FIWNQKRARIAKTILSQKNKDG
	İ	1				GITLPDFKLYYKATVTKTAWY
						QYQNRDIDQWNRTEPSEIIPHV
	ĺ					YNHLIFDKPDKNKK WGKDSLF
						NKWCWENWLAICGKLKLDPFL
i						TPYTKINSRWIKDLNVRPKTIKT
						LEENLGNTIQDIGMGKDFMSKI
1						PKAMATKAKIDKWNLIELKSFC
1						TAKETTISVNRQPTEWEKIFAIC
						LSDKGLISRIYKELKQRHKKKT
						NNPIKKWAKDMNRHFSKEDIY
		1				AANRHMKKCSSSLAIREMPIKT
		1				TMRYHLTPVRMAIIKKSGNNRC
		1				WRGCGEIGTLSHCWWDCNLVQ
		1	I			PLWKAVWRFLKDLELEIPFDPA
		1			l	ISLLG/IYPKDYKSCCYKDTCTQ
L				l		STFTCRQHLPL

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	locatinn of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14590	44958	A	14676	1	2354	MGDFNTLLSALDRSTRQKVNK
1.1270	,					DIOELNSALHOADLIDIYRTLHP
l				l		KSTEYTFFSAPHRTYSKIDHLV
						GSKALLRKCKRTEITNCLSDHS
						AIKLELRIKKLTONHSTTWOLN
				l		NLLLNDYWVHNEMKAEIKMFF
				l		ETNENKDTTYQNLWYVKDTRI
				l		SGMLWYVKAVCRGKFIALNAH
				l		KRKOERSKIDTLTSOLKELEKO
				İ		EQTHSKTSRRQEITKIRAELKEI
						ETOKTLOKINECRSWFFEKINKI
				1		DRPLARLIKKKREKNQIDAIKN
						DKGDIITDPTEIQTTIREYYKHL
				1		YGNKLENLEEMDKFLDTYTLP
				1		RLNQEEVESLNRPITGSEIEAIIN
				1		SLPTKKSPGPDGFTAEFYQRYK
						EELRIKYLGIQLTRDVKDLFKE
						NYKPLLNEIKEDTNKWKNIPCS
						WVGRINIMKMAILPKVIYRFNA
1	l					IPIKLPMTFFTELEKTTLKFIWN
1	ŀ					
		ĺ				QKRARIAKSILSQKNKAGGITLP
		i			İ	DFKLYYKATVTKTAWYWYQN
						RDTDQWNRTEPSEIMLHIYNHL
						IFDKPDKNKQWGKDSLFNKWC
						WENWLAICRKLKLDPFLIPYTK
						INSRWIKDLNVRPKTIKSLEEDL
						GNTIQDIGMGKDFMSKTPKAM
				i		ATKAKIDKWDLIKLKSFCTAKE
				l		TTIRVNRQPTEWEKFFAIYSSDK
ŀ				l		GLISRRYNELKQIYKKKTNSPIK
1				İ		KWTKDMNRHFSKEDIYAAKRH
		Ļ				MKKCSSSLAIREMQIKTTMRYH
14591 14592	44959 44960	A	14677	1 I	3144 2745	
14592	44961	A	14679	1	2742	
14593	44961	A	14680	I I	3264	
14594	44962	A	14681	II	2982	
14595	44964	A	14682	1	3102	
14597	44965	A	14683	1	1443	· · · · · · · · · · · · · · · · · · ·
14597	44966	A	14684	li	3723	
14598	44966	A	14685	2	1841	
14599	44968	A	14686	572	744	DLLTTIRMPTDVKKCSSSLAIRE
14600	44900	I^	14080	1212	/***	MOIKTT\M*YHLTPVRMPIIKKS
				1		GNNRCWRGCGEI
14601	44969	A	14687	1	3588	GINING WROCOEI
14601	44969	A	14688	1	5569	
14602	44970	A	14689	1	3992	
14003	449/1	I ^A	14089	L1.	3774	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ļ.	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14604	44972	Α	14690	755	2415	QELRVSGWSPASAPANT*CPPA
	1	l				OPCPTRRSGGORAPARTRSRPG
		l			\	SRPRTASALPRPAAOKAASRSA
l		l				RASRTNPYTAPKPLPSVESALO
l						EAAIGEAHAQQRVSCTCAAPA
l		l				APPSPAWPWSAARSAAPPPRT*
	1	l				ASAPA*SPL/PVRLKKSPGSSNC
		l				EARGEGWKT/PALPVQLGAHGF
		l				PQPAAVPCASPGSGAGPRWSRS
						ARRTPRGPLORGRTWVSPDPGR
		l				EATPSGTQWPQKTRGSLPGS*C
		l	1			APETPCSAA*SSGTSPPRGAGGC
	1					CPGTVPPALEGHN*PQRWGSHS
	ł	i				AGRVE\TRAEGPQCQQGAPAGC
	1					SLGVGGQRGPPGLPPIPVPGLC*
ŀ						TCRSS*MEAATGSVEKCKRMA
l						SLMQASRWSFLRVAPVSGQVK
						YTPGTARWAEPSSLGSDTTLVP
1		l				GAS/RAPADGGFGLKFLSHLVS
	į.	1		i		DGVLRCGGPRPGTASAPAGLR
		l				AAAA\PSVPSLVGPWRGGLGAG
	l	1				WLGFSLWAFSPTDSLSF*PGPA
		1				LLNLMSALNS*KNFFMVAARS
						CGGGTRGRGGTGPSRGRGRRR
						PATGRK*KKKKKKKKFTSPPEV
		1				PPSPGLPRDNRL
14605	44973	В	14691	1	822	
14606	44974	A	14692	1	2789	MAAVRKQLPRGVLGYGGSGAL
		1				LSTAAADTSLLLGSLQERKSAS
1		l				AGPGYLRSSVAAWSGLWLTGA
		1				SEAHSVVLACFSRKDVEASLSV
		1			1	GLPVHPDASCARRLDAADGVA
		1				GGSPRLPQSTDGWRLTAHSLPQ
	Į.	1				STDGWRLTAHSLPPPTDGWRL
		1				TAHSLPQSTDGWRLTAHSLPPP
		1		1		TDGWRLTTHSLPQSTDGWRLT
1		1	l	I		AHSLPQSTDGWRLTAHSLPQST
		l	l	I		DGWRLTIHSLPQSTDAWRLTA
						HSLPQPMDGWSCRSAQWCV
14607	44975	A	14693	176	440	
14608	44976	Α	14694	114	253	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14609	44977	Α	14695	124	890	GSRAWRDPRLEVAAATTTLFY
l						GYGFVSAELKKASKRMTCHKR
	İ	1				YKIQK\KAPEHHRKIRGEAKKR
				İ		GHNKPRKDPGVPNSAP\FKEAL
				ŀ		L\REAELRKQRLEELKQQQKLD
						RQKELEKKRKLET*SWIFKPSN
				ł		V\DPMEK\EFGLCKTE\NKA\RSG
						KQNSKKLYCQELKKVIEASDV
				l		VLEVLDARDPLGCRCPQV\EEAI
	ŀ			l		V\QSGQKKLVL/IYLNKSDL\VP
						KEDL\ESWLKLF*RKDLPNSGCF
						RASTKTQRVKGKITKVSFIS
14610	44978	Α	14696	553	997	EVTACWQSSQPSLTLGASSAW
l				l		APTLAALEEPFSPPLHCGSPFLG
						WPRLEPAPSAPPWAPVRPEPPQ
						QGPPSAPGRPIPSTTQGLRSADA
						GRRTGTQLHRQPAKVCSF/IS*S
l						QRDHEPTRRNEQLQTRRLKSCN
						THREDLQLHSRRKKL
14611	44979	Α	14697	1410	2036	
14612	44980	Α	14698	3	630	RYKGSPRPHQTQEPSWLHLVDP
		1		!		APGLQVELPASPAPCARIRQPL
		l		l		GGRWDWAPWSRGWCSSGRLG
		l	l			QHRSPWSGWEAQAWRAAGPK
		l				PCPAGRQLRPGEKSSTAPVGRH
		l		l		CWGIQYTLRSHWPGC*VPHSSG
		l				PAGLAGCSGTPAGPQAPHAAP
		l				VPARA/CSLHTSLQAEGVGSSL
		l				GQPRKGLPQCRWGLKGSSNAA
						KVGAQAGEVQARALRTAS
14613	44981	Α	14699	190	383	ASHPLRGLRSGRSLPDERHRLL
						HSAQSHRPPKG*GVRAQGAGL
						T/WQLHL*PQCGIHWVKPAGLL
14614	44982	Α	14700	1	1095	
14615	44983	Α	14701	1	777	
14616	44984	A	14702	1	933	
14617	44985	В	14703	070	996	OLD DOLL COROL DEPONDENT
14618	44986	Α	14704	978	1207	ASLPLRGLLCCPSLPEERRPLLH
				l		GAQSHRPPKG*GVRAHGAGLA/
	l	1		I		WQLHLRPRCGIHWVKPAGLLS
		<u> </u>	1.000		550	LGPHQQSSKSA
14619	44987	A	14705	III	558	

SEQ ID	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	1100	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14620	44988	A	14706	1	1103	I MKPWALKVSVTALKVARLEFV
						PPDVRMCSEFLPCGVKLQTFAV
						SVTALKAARLELFVSPGGLVVS
1					1	LASGVKLQTFPVSVTAHKSSVD
1						PRSEQQQDLLQRAKEQTFHTVE
ļ						GDRSGSPPDSGAQLASPSGSRT
		1				RAAGGAACQSRALRPHSSALG
		l				WSMGLGAVEQGAVLIREARAA
		l				HEPMERVGGSGMAGCRSRALP
						REKAAKARRETEGSAAVSGWL
ł	İ				ļ.	SGIAQWPKIDHEGFRAVKMPQ
		1				AAWLLKAVTFIDLTTLSGDDTS
						SNIQRLCYKAKYPIREDLLKAL
		l				NMHDKVAAGFPAGQTHLKTRL
	1	1				EEIRLAVEDGATEIDVVINRSLV
	1					LTGQWEG/LYSPVVYSALPCNI*
						HSRLFLPSGNALLLWLWCLHS
14621	44989	Α	14707	2	2300	
14622	44990	Α	14708	2	1058	ARSNIRMVTAFRSLPAGSSGAA
1		l				RA\MSAHNRGTELDLSWISKIQ
						VNHPAVL\RRAEQIQARRT\VK
Ì						KDWQAAWLLKAVTFIDLTTLS
	1	1				GDDTSFNI*RLCYKAGYPIREDL
		l				LKALNMHDKGIT\TAAVCV\YP
		1				ARV\CDAVKGTSRPAG/CVNIPV
	1	1				GISRAAGF\PAG\QTH\LKTRLEE
	1	1			1	I\RLAVEDGSLQKIDV\VINRSLG
	İ	1				A*QAQWE\ALYDEIRQF\RKAC
						GEA\HLKTILA\TGELGTLTNVY
	1	1				KASMIAMMAGSDF\IKTSTGKE
		l				TVNATFPVAIVMLRAIRDFFWK
		l				TGNKIGFIPAGGIRSAKDSLAW
		ŀ				LSLVKEELGDEWLKPELFRIGA
		l				STLLSDIERQIYHHVTGRYAAY
		ļ		101	998	HDLPMS
14623	44991	A	14709	576	1998	VLQFLKAACPEFVPSGVPMCSE
		1				YLPSGGFVVSLASGVKLQTFPV
		l		I		SVTALNALRLELFVPPGGLMVS
		l	l			LASG\GSCRSSR*VLQLIKAVWT
	1	l				QRPLGGRWDWAPWSRGRRSFG
1		1		1		EARAAQKPMEGVGGSGMAGC
	111000	ļ	1.0710	-	500	RSRSPAPREGS
14624	44992	Α	14710	2	589	RHHPRHAVQARPFPAGPLAGA
		1				QRGSGAE*RGPQGS\PGPAGLS
1		1				GGESL/PGETDRGADKEERGRQ
1		1		I		RWQRALGRLSAGADPLPAELR
	1	1			I	GRRQRRGR*AGQRAAQEAACE
		1			1	APIQGSGARNGRRG*GRRL*GC
1	1	1		l	1	YQ*V*FPGLTRGWGRGSRPSAV
	1				1	HCGWEPP*AGKPSGQTPRHSG*
	1				L	PAG\VDGLPPKVTGPPPGTPQPR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14625	44993	Α	14711	1	1596	MRLTRCQAALAAAITLNLLVLF
						YVSWLQHQPRNSRARGPRRAS
						AAGPRVSLPPHPLTCVWRCSRP
						AMDRPPAASRPETTVATEFVAL
						VLDGARAEAHALLERMVEALR
						AVSARLVAAPVATANPARCLA
	1					LNVSLREWTARYGAAPAAPRC
						DALDGDAVVLLRARDLFNLSA
						PLARPVGTSLFLQTALRGWAV
	1					QLLDLTFAAARQPPLATAHAR
						WKAEREGRARRSALLRALGIRI
						VSWEGGRLEWFACNKKNQRC
14626	44994	A	14712	225	980	VPSRTWAEYRTRFVPHSLGQPR
		1				SSHIPQSPQTDQTAQGQQALAP
		1				PAGRSGQ*GRPGRRARLGPGER
		1	1			HS*APPPLSQLLLQTFPHLQALP
		1	İ			SAHARPSGQPHADRGHSRSPSG
		1				/DLSTSSVLKVVRPQAGGPKQA
		1				RCPQAGRVCHCPRGGRRGPE/Q
		i i				Q/GG*RPEAPQASPEKGSTGLRP
	i	1			-	SAKDSKNQGTGIFLRCQVAPRP
		ı				EALCCPPPRGADTPVSPCLWVA
	1	1				GAGIPLQPPEGAEGAVLGSLCT
		\perp				LCPGSPDSPPTS
14627	44995	В	14713	405	893	
14628	44996	A	14714	178	499	
14629	44997	A	14715	231	436	
14630	44998	Α	14716	141	393	
14631	44999	С	14717	97	351	
14632	45000	A	14718	119	358	
14633	45001	A	14719	21	614	PPCTSLRPLHAFSGKMTHLNRG
		1				TLSLLDSQLHNLIALQMTCFKD
	İ	1				VEIPNFFWEPSVTPSHRNINMYF
	1	ı		1		PAAVFGFLPISGTLFSYCKIVSSI
	i	1	1	1		LRVSSSGGKYK/AFTTCGSHLS
	1	1		1		VVC*FYGTGFGGYLSSDVSSSP
		ı		1		RKAAVASVMYTVITSMLNPFIY
	1	1		1		SLRNRDIKGVLRQPHGSTVQFQ
		 _		ļ. <u></u>		YLLICSIPFVVWVKKGSKVK
14634	45002	В	14720	162	1263	1

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
14635	45003	Α	14721	696	1791	VWIALKSENITCELLLFKMQTQ
						COVLVYLFPIFOPSFLGKRCPIS
l				l		TDPONLTDVSIFLLLEVSGDPEL
				l		OPSPCWAVPCPCAWVTVLGNL
1		l		l		LIILAISPDSHLHTPMYFFLSNLS
1				i		LA\DIGFTSTTVPK\MIVDI\OSHS
				l		RVISYAGCLTQMSLFAIFGGME
				l		E\DMLLSVMAYDWFVAICHPL
						YHSAIMNLCSVGFLVLLSFFFF\
		l		l		LSLSDSQLHNLIALQMTCFKDV
				l		EIPNFFWEPSQLSHLACCDTFTN
				i		NIIMYFPTAIFGFLPISGTLFSYY
ŀ						KIVSSILRVSSSGGKYKAFSTCG
l				İ		SHLSVVC*FYGRGVGGYLSSDD
l						VSSSPRKGAVASVMYTVVTPM
l						LNPFIYSLRNRDIKSVLRRPTEH
				1		CLISKK
14636	45004	Α	14722	1	1971	MTRKVTLLLRTWRGGCGRLA
						MGMEATPOGSLHPLFQEHWIR
		1		l		AEDFGVLSPMHPNSVQGVDDM
		l		İ		IRLGDLNEAGMVHNLLIRYQQ
		1				HKIYVSLPSPVSTGEPLTGANLL
		i				DREKKMMHRGHCPIALTEGVS
		ł				GPDNPEGRARAKAKAKARAEL
1		1				PLPGAEGTDHLLPSQTYTGSILV
		1				AVNPFQVLPLYTLEQVQLYYSR
		1				HMGELPPHVFAIANNCYFSMK
1		1	1			RNKRDQCCIISPQNRVGAQTDT
		1				VLKVFQHLLISLICKTTLQACCI
		l				PSHPYLSSAPGPPPLTTQCRSYS
ł		Ì				DSLLGPQLFGDVCYNCSHVIEG
				ł		DGEALCSLPHPTCSPGAWVLQL
				l		VSPLNKA WCVSCFSCSTCNSKL
1				l		TLKNKFVEFDMKPVCKRCYEK
İ						FPLELKKRLKKLSELTSRKAQP
i				ĺ		KATDLNSAEGPLAQLGAVGA/V
						TGITHEPHIPQVENGSQQLDDV
		l		l		PALGV\ENWRIFMAERMEAQG
		l				GQRLLLSTMHEEYEFVSPSSVAI
						AELVALFLEGLKERSIFAMALQ
		l		1		DRKATEGGPVGLTKKQGLLAS
			1			ENWTLGQNDRTGKTGLVPWH
		l		1		CLYTIHTVTKHSDSCCTAVPTL
						LEASSGCIFMSYKKHITLTPKSA
l		1	1			DSNKRPSTLGSQFKQSLDQLMK
				l		ILTNCQPYFIRCIKPNEYKKPLG
L		L		L		FSDSFMLIKLIRIDKVQYDSFSYI
		_				

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop cudon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14637	45005	A	14723	11	1197	MNMNIKKIVKOATVLTFTTALL
14037	43003	^	14723	ľ.	1177	AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQONEKY
						QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
						VFALAGSRKTLMTHQSTCPKH
						RASRVGPSNSLQNKQTVKKGD
			ŀ			QTSVRRKRNDLKVEERHKNHH
						LARELTSSAAAASPTGHAAATA
						QAPQRHVEVGRAAAPAGDLGE
						GIVGEQVLGAERLVGPDASRAR
						ALCVLGDRREAGHTAEMTAAP
						PAQVPGPGGPTALRRPSSYLGL
						GVDEQNPMSVDGDEDDVRVQ
						HQELRDCLIKHTILIRGEFVTRS
						LNIAQAADRRDAFVKGIYGALF
						LWIVKKINAAIFTPPAQDPKNV
						RRAI/GLLDIFGFENFENNRIIRT
						LCWRYREARC
14638	45006	С	14724	I	1032	
14639	45007	Α	14725	1	4960	MSTTSIFQQLNLLLSHSHKQKQ
						QIQLPNGRSRYSCLMAAFRMQ
1						YFLSLPMKSLILVGFERLQAIAR
						SQPLARQYQAMRQRTVQLQAL
1						CRGYLVRQQVQAKRRAVVVIQ
						AHGQGMAARRNFQQRKANAP
						LVIPAEGQKSQGALPAKKRRSI
						YDTVTDTEMVEKVFGFLPAMI
1						GGQEGQASPHFEDLESKTQKLL
						EVDLDTVPMAEEPEEDVDGLA
						EYTFPKFAVTYFQKSASHTHIR
1						RPLRYPLLYHEDDTDCLAA
14640	45008	Α	14726	86	370	RTWMTHVWRRRRVQAAPTAY
						SSFRPGTSPCPAQPRAGARA*TR
						SGTSIMPTGFSSHTSCLTVP\PFT
i				I		ASSAGQHRQRHRKTTAAGHPT
				l		TTGAASF
14641	45009	Α	14727	3	455	MOSVLPAHGVSSTRASSRSVTL
1				l		SSPSTSTNCRGRKSSRVT\SPVG
				l		SGPGGGGGGPGSGRFGIDPPRK
		l		1		FAAAGSFSTGAPSSVMVVRVSG
				l		MPSLSW*OYP*SCSTLGCRSRSS
l				l		STISLSSLWA/VQPIAAPPYPELA
l			l	i		TOLPISCR*SPSPCAPS
				L	L	1 APT TOCK OF DE CVE O

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14642	45010	A	14728	338	1782	NQLSK*KAHSIRTTALQOSSGG SQICHOSSADALPSSRCTNIPSST FTPTPLPPRPHPRTCSTLGCRSR SSSTISLSSLWAASR*QPLRTPN STRTQLPRSWPLSTALSPPPPQ ATRLCWWMYSSK*RTWMTHV WRRRRVQAPTAYSSRPGTSPPPCQ ATRLCWWMYSSK*RTWMTHV GRSPATTAYSSRPGTSPPTSP CPAQPRAGARA*TRSGTSIMFT GFSSHTSCLTVPRLHCVLCWTA STSTPQESSPGGWTRPWPSWRRC CTCTGAGRYLQNSTAST/PPVB STYSOTAPSSFEDYRCWPSYHH GSLPPFSVQPG*GCGMSVRAMP SVGPLWSPTRPPQGIKRQNYLL NYWSCWSMTLLAGIIPSST/GS RPADLDYFLGSSQCPRDERTLK GHVSCDPTSISITIGFEKLRQKLE SSSTLADTKSAFLQDSWT/SSLP TLGASARRTTSAS*TDQCPPAP GPLHYSFSLNSTYPSRLCSSTCS TVDSTQIIPMPN*MPFSTCGAFC WCCWFSATTALYFVM
14643	45011	A	14729	148	1040	WCTGSOFSKGTWEILLRKKPSL GNVGALPCVATACLTTVEKRA PRGHPLAPVLGDICPKVOSMSS STSELJCTWONFEFWGRNKYMI WF*WGWALLHOPKMGKGAQG LQEASFYRKDTY*A'ITMP*NPIG ISEFGTU.ALPGPCKNYFERFMS FRELPGKA.AMTDNTNVYVR YKGDYYLCTETNFANIKVDIET LEKTEKMWKPRFRGICPOLQG CSYTEQGFESDDAKSWALQH PKRPAQQDWAMSRONGWAN LGLKIRGEAVTTRAKEVSVUL KYLEVTDGKVWAREKKLRDSR
14644	45012	A	14730	189	507	AAGADTSQGPGGGPPPSQSQR\ PKSPSAGRKGSQLSPQPQKKGL PSPQGTRKSAPSSKATPQASEPV TTQLLGQPPTQEEGSKAQGMR* PPQRLGASTQQTNQEAA
14645	45013	Α	14731	2	356	DGVSLCHPGWSAVARSQLTAT SASWAVCWQGCGETGILLPCW WECEKVQPLWKQYGRLSCEDH LNS*VQVILTTQPPILFPKWLHL FTFPPTGQKDSSFSTTLPTHCPG GRGCSELRSRHCTPAWVTE

SEO ID	SEO ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
i				sequence		
		<u></u>			L	
14646	45014	Α	14732	1	408	FFFFFYTTFFLFFYFYFFEMESH
						SVTQAGVQWYNLSSLQPPPPRF
i						RRFSCLSLPSS*DYRCPPPHPAN
						FLCF**RQGFTLLAKMVSIS*PH
						DPPTSASQSAGHVLTSKWELNI
						EYTWVIKMGTADTGEHKNGEG
		ŀ				WE
14647	45015	С	14733	262	536	
14648	45016	Α	14734	179	745	TDCLHNRCHRSPRSPPPGPPG*R
		ŀ				SRADWWRRGRCWGSWCASSR
						RVC*RCPAPSASFSCCRSPRGRQ
		l				GRTFPPSPCPAPD*TQ\PVSPGDS
						TRPCPPPSSPPAHRAT/GCCGRR
		1				RLPGNASYLASPGKEAGEFPAS
		l				QGDS*PGQYGYIGDSSS/RPTRG
		l				TWMAKASTPRHPIAPOICGA*G
		l				FGEHQEVPVREK
14649	45017	В	14735	1	1164	I GENQE VI VIEK
14650	45018	A	14736	35	542	VRPPSHVTADSGRSPLSLTYLPL
14030	73010	l^	14730	"	342	OEPGDMAAAVPRAAFLSPLLPL
	ļ					LLGFLLLSAPHGGSGLHTKGAL
						PLDTVTFY/KDYGDKLNMELSE
				1		KYKLDKESYPVFYLFRDGDFEN
				1		
					İ	PVPYTGAVKVGAIQRWLKGQG
	İ					V\YLGMPGCLPVYDALPGEF\IR
		Ļ.				ASGVEARQALLKQGQDN
14651	45019	Α	14737	11	908	VRPPSHVTADSGRSPLSLTYLPL
						QEPGDMAAAVPRAAFLSPLLPL
					ĺ	LLGFLLLSAPHGGSG\LHTKGA
						LPLDTVTFYKVIPKSKFVLVKF
						DTQYPYGEKQDEFKRLAENSA
		l				SS\DDLL\VAEVGISDYGDKLNM
						\ELS\EKYKL\DKESYPVF\YLFR
						EGDF\EEPSSHYTGGS*RLGAIQ
	-	l	l			RWLKGGKGVYLGDGLELVCT
		l				V*LTPWSGEFIRA\SGVKARQAL
						LKQGQDNLSSVKETQKKWAEQ
1		[1		YLKIMGKILDQGEDFPASEMTR
	l			1	1	IARLIEKNKMSDGKKEELOKSL
	1					NILTAFOKKG\AEKEEL
14652	45020	Α	14738	3	227	
14653	45021	A	14739	1	424	
14654	45022	A	14740	1	207	
14655	45023	A	14741	2	352	
14656	45024	A	14742	1	441	
		L	1 - // 100	F		

SEQ ID	SEQ ID NO: Mct SEQ ID NO: Nucleotide				Nucleotide location of last Amino acid sequence (X=Unknown,			
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)		
				sequenee				
14657	45025	Α	14743	1	461	MTVDYRKFNQVVTPMAA/AVP		
						DAVSLLEQINTFPGTWYAAIDL		
l		i		l		ANAFFSIPVHEAHQKQFAF\LPQ		
						GYINFPALCHNLIRRELDFFLLL		
	l l					QDITLVHYIDDILLIGSSEQEVV		
	1					NTLDLLIHKRSKEAEHTAASRIR		
						VSCLPEQKSHEQTLPWEQVP		
14658	45026	Α	14744	1	928			
14659	45027	Α	14745	283	1074			
14660	45028	A	14746	450	1970	VRVLSPVEKELKLWKNTHKLL		
				l		SYPTVGAAVTQLQNLTAMGVI		
	1					GSHGARGQVVALNRQRQGDL		
	i	1	1		Į.	QPFTRVTVHWGKGYDRTFWGL		
				1		LDTGSELTLTPGDPKHHCGPPV		
		l				KVGAYGGQVINGVLAQVQITV		
		1		į.		GPQTHPVVISPVPECIIGIDILSS		
		1				WQNPHIGSLTGIMVGKAKWKQ		
İ		l				LELPLPRKIVNQKPYCIPGGTVE		
					l	ISATIKDLKDAGVVIFTTSLFNS		
		1			i	PIWPVQKTDGSWRMTVGYRRL		
		1				NQVVTTPTAAAAPDV/VVSLLE		
						QINTSPGTWYAAIDLANAFFSIP		
	1			1	l	VHKPHQKQFAFSWQGQQYAFT		
				1		VLPQGYINNPALCHNLIQRNLD		
				1		HFPLLQDITLVHYLNDIVLIGSA		
	1	l				IKWVMHSS/DSIIKWKWHVHDR		
	1					AQADPEGTS*PGYGHF\WCPIC		
	1	1		ŀ		QQQRPTLSSQYDTIPWAISQLPG		
1		l		1		DRLIIGPLPLWKGQKFVFTGIDT		
1	1		1	1		YSGYGFAYPACNASAKTTIMES		
						QNTLSTLMVFHTALPLTKALTS		
				1		WLKKCSSGLMLMEFTGLTMFSI		

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SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop cudun, /=possible nucleotide deletion, \=possible nucleotide insertion)
14661	45029	А	14747	1	2589	MGPPYKPPTTWPPQTLSYHSTT PVKLQSLSLPPTAARGATEPCH LQAPASSVQQMTPPAPGLKHL
						GGKSQMCLEQDGRSGSVRGSP CHAGHWVAGAGFGKGTTRTW
		l				GHVNLIVPPNCPTASSSAARDS
1		l	1			APNOPLLGVVTAQIVPPTCPPA
			ŀ			MTVQELIAPLTLPPAAGCRRGV
						RPFQTRVWLLAQLLRARLGGP
		1				GEGVVTCHSWRLERVAPRLAS
		1				LWWALAAETAARLDWAAKRP
1		l				SAWGSIKAAEFSSKATPRLEGA
i		l				SSRTLWVSYIQAKETTTGKNGP
						VEPSHLWEAYCRATAPHPAVH
						SPSAEVTYTCGIVNLLLQGLVP
						KSHLETHTGCHVVVELIICSVV
						LPLISRLSDPDWIHLVLMVIFSK
		1				ARDPTPCPASGPEQPSVPTSLPL
1						IAEVQQLPEGRAPSPVAAPGFLS
1						KQIQDALCALEGSQALEPKGGE GSEGVEAEGPGTETETVLPVST
1						LNFCPEIQIDIADKEIEQGEVTTS
		1				VTALLEGLEKTCSSRPSCLEKD
						LTNDVTYLDPTIFSFEPLSSPDG
		l				PVVIQNLHITGTITAQELSGTGL
		1				OPCTRYTVKCETSLDGENSSLO
		1				OLAYHTVNHRYWEFLNLO/SH
	ŀ					QEKKPDL*KFIKN\IEKMVV\$AI
						VDNLKTAFPCSESQGPTEELSE
		l				AETESKSQTEGKKARKSRLRFS
						SSKISPALSVTEAQDTILYCLQE
						GNVESKTLSMSGMESFIEKQTK
14662	45030	Α	14748	478	3507	SKILRAVIWGSHLVRSAMKTET
						VPPFQETPAGSSCHLNNLLSSR
		1				KLMAVGVLLGWLLVIHLLVNV
		i				WLLCLLSALLVVLGGWLGSSL
	ł	l				AGVASGRLHLERFIPLATCPPCP
	l					EAERQLEREINRTIQMIIRDFVL
				i	1	SWYRSVSQEPAFEEEMEAAMK
		1				GLVQELRRRMSVMDSHAVAQS
		l		l		VLTLCGCHLQSYIQAKEATAGK
		l				NGPVEPSHLWEAYCRATAPHP
		1	1	I		AVHSPSAEVTYTRGVVNLLLQ
14662	45021	<u> </u>	14740	ļ. —	412	GLVPKPHLETRTGRHVV
14663	45031 45032	A	14749	34	417 350	BEDCOECCUA CDVEMEVEVIVE
14004	43032	l ^A	14/30	34	الادوا	RERGQEGGVAGRVEMDVSVVE RGAYADOIWGRPKGGESRGEK
		1		1		RKGRWVQRVRALRARVTPAGL
1			1			GECKWGRGRGD*VRRVKPRAL
		1				RERWGDGSTVSDSTPMKPGLE
	L	Щ.	L		L	KEK WODGOT VODOT FINKPULE

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	coden for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14665	45033	A	14751	ր	1604	MASKRCIRDFRSRGGKLKSFQL
	l .					YATFRKEGCGPVLANGNPGGG
	i		1			RGQEVSATQDLRRGWGIASSAS
		l				RFPWRQSWPGRPRLGNYDSGA
						PGEAKMPSKGKDKKKGKSKGK
		l			}	DTKKLIKTDESVVDRAKANASL
		l				WEARLEVTELSRIKYRDTSRIL
						AKSNEDLKKKQCKMEKDIMSV
				İ		LSYLKKQDQEKDNMIEKLKQQ
	ı	l		l		LNETKEKAQEEKDKLEQKYTR
						QINELEGQFHQKAKEIGMIHTE
		l				LKAVRQFQKRKIQVERELDDEI
	i					NDLLVKEKIMQLVQQRSQIQTL
						QKKVVNLETALSYMTKEFESE
						VLKLQQHAMIENQAGQVEIDK
	ł					LQHLLQMKDREMNRVKKLAK
	1					NILDERTEVERFFLDALHQVKQ
		l				QILISRKHYKQIAQAAFNLKMR\
		i				EHVQEEQNIPKSEHLMAESTAPI
		1				V*IRIFWRPKNGHILKEMWILEI
						*PGSRRKKYCDCSLQK*MAVLL
	i	l				GNTTQSSRPSSSRTMLFLDSGET
				l		KEFGDESKLQDKIFITQQIAISD
		l		l		VFW*SGATHYSKRTSGV*HSGK
		l		}		SESLQPRGQRLI
14666	45034	Α	14752	95	473	
14667	45035	Α	14753	168	449	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14668	45036	A	14754	li .	2000	MAFEOAKOVVKLALDLWPIGD
14000	43030	<u> </u> ^_	17737	l'	2000	GPVELQVTVLDQHANWSLRQE
	1					OHGKRGPLOFTDVAIEFSLEEW
						HCLDTAORNLYRNVMLENYRH
						LVFLGIVLSKPELITCLEOGKKP
						LTVKRHEMIAKPPGQKNWEKQ
1						ALPLTGCLTKYPSWAKNIDIIVT
						CSHFARDLWPEOSIKDSFOKVT
						LRRYENYGHDNLQFKKGCESV
]		DECKVHKRGYNGLNQYLTTTQ
						SKIFQCDKYVKVIHKFSNSNRH
				1		KIRHTGKKPFKCIECGKAFNQS
	Ì					STLTTHKKIHTGEKPFKCEECG
				ŀ		KAFNWSSHLTTHKRIHTGEKRY
l						KCEDCGKAFSRFSYLTAHKIIHS
						GEKPYKCEECGKAFKRSSNLTT
				i		HKIIHTGEKPYKCEECGKAFKR
		i		l		SSILTAHKIIHSGEKPYKCEECG
						KAFKHPSVLTTHKRIHTGEKPY
						KCEECGRAFKYFSSLTTHKIIHS
						GEKPYKCEECGKAFNWSSHLT
1				l		THKRIHTGEKPYKCEECG\KGF
1				i		KYSSSLTTHKIIHTGQQPFKCEE
					l	CGKAFKCFSILTTHKRIHTGEKP
						YKCEECGKAFNSSSHLTAHKRI
ŀ						HTGEKPYKCERCGKAFKRSFIL
						TRHKRIHTGEKPYKCEECGKGF
						KCPSTLTTHKTL\EIHAGKKPNK
1						*E/ECSKAFKDTSALTRHKIIRTG
						EKPCEFDECRKAFNQLSTFTKY
14669	45037	В	14755	269	1131	
14670	45038	Α	14756	2	149	ALFRAGRQECLSLLKLCPQPPL
						SPTCFVPGR*GFYLHFFQRCAA
				l		QRGGI
14671	45039	Α	14757	2	175	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop cudon, /=possible nucleotide
NO.	sequence	1100	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14672	45040	A	14758	1523	2667	PLLVIHRQTGSGEDLQQTPADL
				l .		QQRDLLERKLTNRNE*HQHQQ
	1			l .		KRHPLKNPTQMSP/CIKDQRYIN
						PRSTIKLELGIKKLTENCTTTWK
				1		LNNLLLNDCWEKQERSKTDTL
ļ		1		1		TSQLKELEKPEKTNSKASRRQEI
ŀ	1				į .	TKIRAELKEIGTRKTLQKINKCG
1		İ				SWFFEKINKIDRSLARLIKKKRE
l				ı	ŀ	KNQIDTIKNDEADITTDPMEIQT
						TNREYYKHLYANKLENLEEMD
						KFLDTYTLPRLNQEEVESLNRPI
İ						TSSEIEAVINSLPT/KKRSPGSDR
				1	l	FTAEFYQRYKEEL\VPFLLKLFQ
				1		TIEKEGLLPNSFYEDSIILIPKPG
					l	RDTTKKENFRPISLMNIDAKILN
				1		KILANRIQQHIKKLIHHNQIDFIP
		1		1		GMQG\WFNIHKSINVIRHLNR
14673	45041	Α	14759	640	1771	APVARMTA/PLFYTALLVFSAL
1	1					GNILALCLTCQ\KRRKINCTGIY
1		1				LVHLAVSDLLFTVALPGRVVC
	ļ					YVLGSSWPFGKGLCRLTAFVL
		1				YTDTYGGVYLMACVSVDHYPT
					ļ.	\VVCAHWGPCLRTAGRARLVC
	1	1				VAIWTLVLLQTMPLLLMPMTK
						PLVGKLACMEYSSMESVLGLPL
		1				MVLVAFAIGFCGPVGIILSCYM
	ł					KITWKLCSTAGRTQ*PAGKDTT
	ŀ					GGASPGGPSDQPEKTPPAGAAQ
						EDPVTPRERTPPAGQPERTQ*PA
				I		GKDATGGTAREDPVTSRKGRH
				l		WRGCLLTLLMLVAVVVCFSPY
						HLNIKQFMARGMLHLPSCAER
					i	RAFLLSLQATVALMNMNCGI/D
		1		ł		PHYFFASTHYRKWLLGILKLKG
		L_				SSSSSSSSTLY_
14674	45042	Α	14760	I	401	MVLVQWLTAITSAFWETQNKG
		1		1		SLQFEDK WDFMRPIVLKLLRQE
		1				SVTKQQWFDLFSLGATLQDSSG
						PLLTGKLKKKRKVSGTKY/IEAS
		1				SATSLKATTDTHHLLLYNPFWI
			l			PLTLI*HLCWLQRFA*SCDTDPH
						P
14675	45043	Α	14761	184	2720	
14676	45044	Α	14762	34	251	
14677	45045	Α	14763	2	452	LQIEII*ALEELAAKEKANEDAV
	1	1		1		PVCLAADFPSAGMGSSCDGQE
	1	1		1		DELDIKSRAAYNVTLL/NFMDP
	1			l		PKIPHLKEKPYFGMRKMAVRW
1	İ	1				HHAENLVDRPQ\EESEDDSHLE
		1	l	l		GRDLDTWHVGVKISWDVKTPG
L		L_				LATPQGDCYFMLGNLWRTKIIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14678	45046	Α	14764	1	1701	MPFKCKKQHTFNKTSPKEGAII
1		l				FTHGCLCTSPFNSELSEHLGNL
						KLSSWEHLHHRSQQTLSVRLH
i	l					VPTPQPVEHEFIQLIYIHTGAKL
			ĺ			QASKVIIATKDGKRRKSPLWQL
						KYPKLILREASSVSEELHKEVQ
						EAFLTLHKHGCLFRDLVRIQGK
	ł					DLLTPVSRILIGNPGCTYKYLNT
						QLFTGP\FPVKGSNIKHTEA*I\A
						AVCETFLKLNDYLQIETIQALEE
						LAAKEKANEDAVPLCMSADFP
1						RVGMGSSYNGQDEVDIKSRAA
						YNVTLLNFMDPQKMPYLKEEP
-		1		İ		YFGMGKMAVS\WHHDENLVD
		l				RIQLQLWDPVDGTTASSAVPKQ
						GRDKRREVPNTLSQPNSQEGLI
1	i	1				SAFHSRCVHAVFYKLTSAPKTT
		ŀ		l		SLPDVPSDLLAEGLAASRAGQA
	İ		ĺ			LWRCDKKSLLTLVRKINLQGK
	İ					RDGGRSATNQQSQPHLDNTGL
	1	l				ERQLEQK VGRCRSPWRGLGHL
		l				EGNAQKGHLLHGVNKNGQHL
						LRRHGVSIRAGRAPHLAAADT
1	j	1	İ			APGRISCLGHSEALKPRIQAGGL
1	1					GAESIWQVLWEGGSNSSVFCR
						HDEEPAAELPLPGKQASGGHM
14679	45047	В	14765	16	1462	
14680	45048	Α	14766	89	566	MEQGSIITQAWREDALVLTKQ
1	1	1				GLVSK\SSPKRPHVCHILK\APFC
						CSCTQHVGQSSSPR*ANPHIRRE
1						STPIAMSDQLQRLQYQFYQIPG
		l				TCLLPEVTEKNQGRICMVTDLD
1	ı					ETLVHSSFKPISNADCLVPVELE
						GTMHQIHVLMRPYMDEFLT*M
L						EEMF
14681	45049	Α	14767	1	305	LSGLEMFQAGAGAPSSPGY*GS
1		1	1	1		CWAISWSGPPDKHSNRGASCG
						LETWAGAAATDTAGPHPAAGR
						C*A*AGSGGDQGTGSTAPLPAT
I			1	1		P*PLQSLV*AAALPA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X≈Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14682	45050	A	14768	1070	2653	SLKPCMACL*HS*EKTLSDSSG
14082	45050	^	14/08	1070	2033	RSWICAVSF\GSPGPAAAPLPEL
						SFPLRPGYTKLLPHATEEVSPFG
						SGAGRKLGPMLPSTAELCPGTR
	1					ARWSEGYRDLGRQAQDLQAG
	İ					RKSVVTWALEGHPREEERAAA
	1					KVQQCQRMWEGCSDPQELIPR PGRNPDPVPDSAVALLPWGPSR
						VPDLGPDSEHR/PAHSRVOYWL
					l	
ļ						HKAPAPPSPSPQLAEPATPPTRP SSPGSRSPSSWHRLRSWSLNAD
						PW/RPRPADLPPGNPAPPSRGGA
1	ŀ					HGTWGKDVRPLGRGSRLFSVV
		1				
		1				CGRREVQRAARRVGTTGPGWR
	1	1				APGREREAASR*QPPGTG/PASP MGRASSSGSRVSPGPIRRTAPW
1						EGCSCRSRHHWGKGLGPPDGE
l	1	ŀ				PG\PGSS*VAQPTVGGRRSLGPG
	İ	Į.				GAAPARDAAAATPPGDAAARE
1		1	ł			GSGRHLRAPASRPQHQPRSPRS
1	i	1				HPPASATTASLQKFRSASSEPPS
	l	1				EDRPFDFSPTAPPOCPRVAPSAG
ł		1				AR*R/AGGRLPE/PR/RPGPWGG
		1				GSRTNLTIPTSLHGPTPAAARTS
		1				PHRKSR
14683	45051	A	14769	748	1073	SQHFGRPRWVDRLRSGVRDQP
14005	43031	l^	14707	/40	[1075	GOHDKTPSLLEMOK\IAGRAGG
		1				CL*SQLLRRLRQENPLNPGGGG
ì		1				CSELRSGLCTSVWATGRDSI*K
	١.	1]			KKKKKKKRKPWSLLSLIVLVOF
14684	45052	В	14770	59	466	Madadada Wasasarrar Qi
14685	45053	Ā	14771	2	3184	HRTGIPGSTISSPGGETVPE\MTA
1.000	1.5055	ļ.,		ľ.		AMRERFORFLHEKNCMSDLLA
1		1				KLEAKTGVNRSFIAL\GVIGLVA
		1				LVLVV\GYG\ASLPLQPLLGFCY
		1				PA\YISIKAIGE/SPNKEDDTOW/
		1			1	V*TYWVV\YGVFS\IAE\FFSDI\F
		1				LSW\FPF*LHG*KCGFLVGWCN
					1	GPEAPSKWGLKLL*QAASSGPF
		1		ľ		LSLEATKSQIGQVLVKD\LKDQ
	İ	1				GPKETADAIH*KKPKKSLP*ILL
		1				G*KKKK
14686	45054	A	14772	14	455	POFPCRRFRALSEEELDNEDYY
1.7000	1.505.	1	1	1	1	SLLNVRREASSEELKAAYRRLC
		1	1			MLYHPDKHRVPELKSQAERLF
	1	1	1			NLVHQAYE\VVERRRTPAEIRE
			1			EFERLOREREERRLOORTNPKG
1		1	1	1		TISVGVDATDLFDRYDEEYEDV
		1				SGSSFPQIEINKMHISQ
14687	45055	A	14773	328	479	STATE OF THE STATE
.4007	1.5055	1/1	1. 1113	P-0	I	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14688	45056	A	14774	15	1095	IKVAKMATALSEEELLÖNEDVYS LLNVRREASSEELKAAYRRLC MLYHPDKHRDPELKSQAERLF NLWHQAYEVLSDPQTRAIYDIY GKRGLEMEGWEVVERRRTPA; EIREEFERLLEREERRLQQRTN PKGTISVGVDATDLFDRYDEEY EDVSGSSFPQDINKMHISGSAE PLTATDTAILSGSLSTQNGNGG GSINFALRRVTSAKGWGELEFG AGDLQGPLFGLKLFRNLTFEC YTTNCALOFSSRGIRFGLTTVL ARNLOKNTVGYLQWHCSSPLL VQVRPHRNTRACAPEPSFRFFL HVPTWDAECSGARTPSTAWTS AAVKLREACLSGPGSGSHQLLL LTPRSKRRTGGG
14689	45057	В	14775	508	1765	
14690	45058	Α	14776		639	GRRRHRRHRRCSRGCYLPT CTNTQQPLPPREPLELPLARTP DLSMQNSEGGADSPASVALRPS AAAPPVPASPQRVLVQAASSNP KGAQMQPBISLPRVQQVVQQVQ PVGHVYPAQVQVVEGGDAVY TNGAIRTAYTNJEPGMYAPSS TASYFEAPGGAQVTVAAS/SPP AVPFHSMYGITMDVGGSPHVFS AGYLHIGGMDRFELLANL

1848

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14691	45059	Α	14777	1	2091	MKPDDNSTQNEGGQSFEERHE
,		1		ľ		TLQRVKFRGGPKTAPGRKLIAF
						LGOMNSGTSGGORVPGEEELM
						AGGATCLPNTEAHQPIRMKNL
						HTPAFRLCPLTLQDNEADAFW
				l		HQAFVDWGMPTGKKPQTLGLT
				ł		YKTTHDLSMONS/EGGADSPAS
		1		i		VALRPSAA/APPVPASPOMLPLE
i				ł		LVILGVV/PLDKCIPAQWSKEG
				l		QTNSYTYNPEPQMYAPSST/AS
				İ		YFEAPGGAQ\MVGITMDV/GGS
						PIVSSAGAYLIHGG/MDSTRHSL
						AHTSRSSPARSFRSNSKAPSSRK
				İ		FCC*LPTLSSKTL/SLFSKNLLRK
						LPLGKHSPDSVQWPWIYGLLV
						AGSESLASLLFPKASSSGETLGF
						ATGQLAGESRLPSLLAFILVPID
						VAWGKEQLVTQTTVYRVQGS
				İ		GOHLNKAWQEPALVNICGMGE
				I		KVDVOVGLCOHRAHLDGWNS
				l		SRAASVWPGETLTHHQDQCQA
		1				NLPPGIETSILGTKGARYPSPQE
		1				QGGGRVWLERLDPRG/AVSGT
						DL*CLPQSDVPPGRA*SGGTGD
		l		ł		KSISTHDKSGRGNSKYHY/YGIR
	i					LKPDSPLNRLQED/TQYMAMR
İ				1	i	QQPMHQKPRWCRAELGGAGSP
				1		L\GMEDVSHVFPEFPAPD/LGSF
				l		LLQDGVTLHDVKAL/QLVYRR
			ŀ			HC/EDVV/MNLQFHYIEKLWLS
	ĺ					FWN/SKASSSDGPTSLPASDEDP
						EGAVLP/KDKLISLCQCDPILRW
14692	45060	Α	14778	94	3006	RTRSLTRKAMAEHAPRRCCLG
				ĺ		WDFSTQQVKVVAVDAELNVFY
				}		EESVHFDRDLPEFGHVLDVHG
						VHVHKDGLTVTSPVLMWVQA
						LDIILEKMKASGFEFSQVLALSG
			Ì			AGQQHGSIYWKAGAQQALTSL
1	l					SPDLRLHQQLQDCFSISDCPVW
			Ì	1		MDSSTTAQCRQLEAAVGGAQA
						LSCLTGSRAYEFNLVCDRKHLK
1	l		1	ŀ		DTTQSVFMAGLLVGTLMFGPL
1			1			CDRIGRKATILAQLLLFTLIGLA
1	l			1		TAFVPSFELYMALRFA\GL
$\overline{}$						

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14693	45061	A	14779	1	710	MGPAAGAPTSRASRGPRAPALR AAVGRRCPRPEMPCQCHPLAR CGLDPAWTQLPQETAGAPRQD PRCTNQLICEHLPLEGLGFGYL GNKDSLRHVKVVAVDAELNVF YEESVHFDRDLPEFGGLASSAV TQSSS/DDLQQVP*SLQSLHAA*
						SLQTNASPFTPEHSFLGKRHPK GAGQKRAESPTRPPLRLQEAAP GELQRPLPYGTQVPAGLCPSVS GLRLRARALLPYRTHSLSV
14694	45062	В	14780	1	252	
14695	45063	Α	14781	73	198	
14696	45064	A	14782	762	1080	AGRAEMQQERESHEEAKDPSP LSPREGSQKQQQNVNTSTPSHC PR/PPLRMPPPAPRRPVP/CDHPR AEECERKAQHWQAAPPAAPVR DPLGEASWAPESGGDVESLYI
14697	45065	A	14783		309	MTKTEREONEOLDGGGYGERL GEYSLFLLCVPSELNOGKCERV SSYFGMNLSGVLKKLPFPPTRE SWPPPHETGTPINHHGGVSNG ALGCAQRCSLHIIATNAGGWER GGVSNSRLFFFLLSASFSDAKIK PAPSPIDHLRAEEC*/PPPLSQPP ALVAWCREHLWAHPKAPLLTP
14698	45066	В	14784	1	471	
14699	45067	В	14785	1	1143	
14700	45068	A	14786	1992	2205	WRSMSGLRKRQLASQISYCPC WS\CP*EESHCRRVSPAPLA*LL SPCPAPQQRSHTDPPFFFKHFP LPFSG
14701	45069	В	14787	389	973	
14702	45070	A	14788	287	525	TNPFNLLQH*KKP*CYTYISITD PTDACHKNKAFSSPAVSQLT/CI FSSTAGSFFVLSGHSCGHLAVFS LWPRSPQLPGTR
14703	45071	A	14789	475	616	
14704	45072	A	14790		704	MTKWKSNPQIIMELQRTLSNP NNLEKSNVKTYYKAKIFDDNLI QLTFQQHKQDRDTEQEKKEE EREKRKQQQFPLRKKAEM RCFGFLTGPGPNLTPHGKTWSR LPPKNHRRPYTSRPEGPRVKA* NHCPFGRRGTRLPPEAAKRAQ AEEAAQ* WPTRSTLRLEVTPRA MPRSPSEDPRVAGVHPAPPRGP RSDS/APLRPQLRPPRSIFSLASE PAAAGNARGRHSVLALY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14705	45073	Α	14791	200	408	SPHLMSINLITFANFLFPCE*RNS
						QFLKS/SIFVMWMIHNGMKCRQ
		l				LDTCHCRHGTCKDSICPGLEES
				l		PLP
14706	45074	Α	14792	240	771	GRGMQSPAQVDPAAKRGNIAG
1		l				HTAHLWPKGGTLPLCF/YHLLI
1						MQIREQP*SSLEYPTF/RILSLVS
1		l				GTKLNKILANRIRQHIKKLIHHN
ł						QVGFIPGMQGGLPTPGSEQGHL
Ì		l				RVSPSSPSSTVTLKLFNFMEPFE
					ĺ	NLNLSKGKASVHTITQENTDSL
		1				KVQGSQFSQGHEKRTLDLKVS
14707	45075	В	14793	I	1557	
14708	45076	Α	14794	150	1668	ESDARQGPFPLENTRPHPSCMQ
l		l				NHGICRSLHPGLCGCPPTARDA
		l				APATAGGVPAY/GPLTGAPPG*
		l				LPGAGIGSKQSSTMGPSGAGSC
						LGL*APSQPAD/PGSIQGVLVIA
		l				DTE*PLVGGDGGLEAEHQWQ*
		l				EAGP/GGLIPKHMNPQSRQQSLF
		l				GRRGYRDSSLRSVLGRTGQQT*
		l				HAELGAGVMGPGGNGR*GH/G
		l				PRGLPCLVQGRRGPGVAAVAA
ŀ		l				GVSVAVPSGMLSENGPGCSK/P
l	1	l		l		HSGDAGQRRGLRENL*GVRALI
		l				WGAASSFLLTEQDLQELADCL
İ		l		l		HSGRKTCFFYFQGRGDVAPGPV
		l		l		APAHHRLAGISTVVPEGPGLCV
		l				GEDRQLPG/GNQRAEPPP*LVRS
1		l				VSSLTGLPALGPRQMPNAGAG
		l				NGSGK*RGVRGSGPDWDCTGT
		l				HGGAGTGKLLWAGLSLFPLSG
		l				KSEAHQQRAAGHGYGRGLGSP
		l				GRPPRSRKGRVLGSESAPA*GSP
ł		l	1			QPLRSARQKPESSCPRQGQPVG
						RGPGAPLEAGHCSFPRPLGPAR
14709	45077	A	14795	3	374	SLHPDFPQGLQA RRPPSWG*RGCSQLPECAADGS
14/09	430//	A	14/93	ľ	3/4	LEDHGPLPPMVSCFSRHPAGRF
		1	1			LRVFPPSPLLAVDCCWHNNSYV
		1		1		
		1				VAVDCCSATSCSLLLYPSQEFPP
		1		1		HKFLVYLVLSWHQLLGGLELIC
L	L	L	L	L	L	TQQPVNPVVSDG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first eodon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14710	45078	A	14796	73	890	MSILVRTDGKVSFPHSRAASSPP PRFAGSLPWTPKGHGFTSFPRL EGYCPS/REGPSDWGRNSFRVSS WAWRPSQGVSHHTFGQGP/PSP GNDDK*VATRKSPGGPGSSPP GWSLQSTSSPGATRTKGNLNTL KG*KQVVD*WVGVGTGWEGS GAPQSPQPSREWBLSPRGVGVV KETWPSPWSGR*APPCPGQPGG QQTHMERGGRG*G*GYGQGG GP*GQWSVTNGKDTRAWSPPP RQRNGRWPAPPPFTFHYYYYY YYYYYYYYFFFG
14711	45079	A	14797	273	678	TEESTSVLSVPNQTGAGAARIGI VSLRLCGLYVTSPPRAARMHIR PHOPWPGSSHHENGEAGPESÆ GAPASASGRGAPEGQAWPPRD AVLSSLMLALFRTKSARHLVLG TRQTAASPFPFFSQPSWYCSLG VDGG
14712	45080	A	14798	551	743	FIRITISAGSSSWGSCRPRAPAP FS WSRPCLQGPRG*CRGLHSRF YAAP'VQGHPVHILPYSKFIRRG RAALRPRQQWGLDSVLGTAVC GGGPRSESDIGSGR*GPEFWYIS PPVPPHVHRGFQQLINCARQVG KGGGGPGCPMLGNHSSQVGL GLGQDQDSLESPGRVSPSGP SAASDPGG*GFQGNPDPVHPAP APPDW

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14713	45081	A	14799	705	2299	KREWRPRHHPLGPPPRCAGRSA PLSGPASPFS*WLPPGHGPCGR MCSRAARGGDVTYRTHNTRAP
1				ľ		SCPSLHWVGERG*DAHTRTPSC
						PSLHWVGERG*DAHTRTPSCPS
						LHWVGERG*/GCPHSGLQLSFP
		1				PRSAGCCKALFEAEKSKGDTGL
		1				GKRGWAGGGAGGAARPCICNP
						TVPCPQSTQSPPV/PPGTR*QLG
1						SSASHSRAASSPPPRFAGSLPWT
		ĺ				PKGHGFTSFPR/RGGVLPFPEGP
		1		1		SDWGRNSFR\GPAGPSMW*PAP
		1				PSDAQVGHGPGRAGLQRCRGR
1						GGHTPFVLAAAAIDSAYFS**/D
ı			İ			P*VFGSNFPHLSCFPG\GCGEGN
						Q*ARRRPLAGAAALPGPNLES
						WVCELSVAGPAVWDPVPSP/GD
1						PTTGPESP/RNSAGACGAQGSM
						R*T*GTPATWSL/GPPPELPGSR
1						HQRQCECGPAHVQRERMPGW
						ACMLKSVAQRGFQRR*QLSKA
						QAAAWRV\PGGR\GAGKD*GW
						RRWGVSRGGGGLCCARGGGGS
						SVRRGEVEGS/GCARGGGGCAG
						WRALSPSSLPISGAQELGS
14714	45082	Α	14800	1	267	CSAGGPWRAPQPRRFHRRRRP
						AQLPPPLPLPPLPASPRIHNRTF\
ì						PRPSQRTPPPAALGCPEPGS/RS
						QGRGHARPPGSGEGDPTVSSPG
14715	45083	Α	14801	218	430	
14716	45084	A	14802	9	201	
14717	45085	Α	14803	1453	1936	EVEKHLCQG*ELLRAQHN*AA
						ACRRPRPPAPGPQCSAGGPMAR
						APAPQVPPPPPPCSA\PPPLPPP
İ				i		LPAS/HAHPQPHFRHGRRSALLP
				l		RPPWAVRSRGALAGPRTRAAA
					İ	GLRGGAGAAPAPADARFPASSP
			1		1	AE*PKFPQNSARALTGFPRCTD
14718	45086	A	14804	1	580	PTVSSPGY TSGCSHSLHKPAGVSTRPPTVA
14/18	43080	^	14804	,	1380	SICAPRPRTSPLRRGSLTLWPFP
1						GGHLTLLETP\GFDSCPEEVEGP
						RFSTFKTS*ARGGRRQRCAPRG
						RREVGPPSAAESGRARGEPPRC
						RLLCRGGK/DPQPRAGEGEAAA
1						GTPEGSRAGASGSEGALPSAAS
						PGNSSGRSRPEAGANPKGAPRR
		1				CVWGEPGGRRTRLFPDE
	L			L	L	C * #GEFGGKK IKEFFDE

			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	detection, (-possible nucleotide insertion)
14719	45087	Α	14805	279	1148	ALHEFAMSRSLHASLH*S**STC
						P*QQPAPAPAGEPRARFPGKSSS
				ĺ		ASSGPG/WPRNHGKQLHLPAQP
						Q/QQGRISSPRVAGGSGCTGRA
						LLLAAGLQNSTRSLNPRDPPTIL
						SHRPLQGM/PDAP/RGCSHSLHK
						P/SRS/TPLSGQL*PPSAPHAPAQ
						APSVAGPLPCGLSPGDT*RYWR
						LLRFDSCPEEVEGPRFSTFKTS*
						ARGGWRQSRVWASARGASTL
				1		QASVQGR*NPQPRAGEGEAAA
						GTPEGSRAGASGSEGALPSAAS
						PGNSSGRSRPEAGANPKGAPRR
				l		FLGGCLL
14720	45088	Α	14806	765	4270	SQDTPGDLQCHGDLL*DGAVLS
						EAPGGERVPRDD*GGVCPEGVP
						HVSLLRDSVAEALQPRGEAGG
			İ			DQVPEAHARPPSAQR*PAGAGL
						RLHPPAAGGVPGQSGGHLPKIQ
						DDLTSKS*TNSITKTLFPNKVTL
				ĺ		SDSGRA*ILEGHRSTQGSLGLEA
						DPGTVSHHQHPCPPNAATHHF
						HRTARPGVQQGPGPASVQQPES
						DGDGALLRSPCSLLPQGADEVL
						LQPDGDKQGGRPRGDSESD*G
						YSERR*AQDEYQGH
14721	45089	В	14807	133	2943	
14722	45090	Α	14808	1	814	MEEELLRELLYSENWWSIPYND
						TIQELCCWAVLKRTFINGNTSR
						KPGMKMVTGLKKQERRSKQW
						GCEKEAKFVPDMWIVLEVGRP
			l			EERGIDGKRWGKSCSVEQSVW
			ŀ			AFQNPIWSKAQTLLNSMK/AYE
						KGKEGVEEKFEARRSWFMRKK
						AISHLHNMEMQGEAASADVEA
						AVSYPEDLAKIIDESDYTKQWIS
						NVDKTVFYWKKMPSRTFIATE
						EKSMPGFKASKDSLTPLLGTNA
						AGGF*LKPLLTYHSQTSRALKN
						YAKLTLLMLYKWNK
14723	45091	A	14809	123	515	PKQSAKTIWTSTARPEDVVTCS
			ĺ			FKEEFLPGNTHHDFSIKNRSDPE
						LEVSRPLARPENFPANRSHSLLE
			İ			L/RPCRGRWAAHWTGCSSLLD
						ASSAPRTDQPSLWRNLPTASLG
L		١.		L		GPWSHNQPGIVFQNTYLMKWP

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14724	45092	Α	14810	279	1136	VDVDLACVRGLLCSCDEFLSW
		l	ŀ			LALNRTVLATPALPSCLSQMAS
		l				SLPGLCARTRRGERADSQRGTC
		l				AFFSPARTASQAPLVASSVRKQ
		l		i		RVWAPNPLNCHTLPPSRGSQRG
		1				GYIGEEGVFGSRVVGIWVPGPN
		1		1		GSPWPLGVQR/PECVLLGVGSG
		1				S/SSHELFSEADFISRLLAPSPSP
1		1				APAS/MGNAIEEEAPELGPLGRV
		1				FVDDFHYDYNFIHLHEDLCYGP
		1		Ì		LEGPDLDLAGTGDRTPPLHSRP
		1	l			RASLSPSALRCP*ALGF/WSTLA
						WGSPNSSSRVPEAQPPAPSLAE
14725	45093	Α	14811	1	1597	GGRKAAVGVETRPQGSAQPTW
						LPVHGDTCTTLLLPCRAWWRL
						GGLPWAASVVGAKCVWTPPER
		1				GLLMLHKPMPGWSSGYRVVLP
						AVAFFRLRAQVTKLGEEVSLRF
		l	l			LKREAKLCGFLQKSFLALEKR
						MKASESSRLKLEGSLRGELESR
						WEKLRGLMEERLRALQGQHEE
						SHLLEQCQGLDAAVAQLTKFV
		l				QQNQASLNRVLLAEEKAWDA
		ĺ				KGRLEESRAGELAAYVQENLE
		ļ				AAQLAGELARQEMHGELVLLR
14726	45094	Α	14812	150	1625	CLGKAFSCRELNAYLRLPTFFIV
						FSLRQPSLRMHSRAWVKTCGA
						MPCSCCRIRHCSSVTDMAVRSL
			ŀ			ALLQLVNRTPLRRRQISAACQR MSTFLFFRACTF*PRNRSSSGRH
		l				/DLTARPRQAILRQNFLK/RAQS
		l				RNLTAPNSPR**LLVKTYRQTS
		l				RPVSAAH*CPR*RGRAALAAAC
1		l				HSGSPGARRQTAGPAEGPAR*T
		l				RRDFL**RAACLRRRWKWPE*L
		1	ŀ		1	*RSRSSSP*SOGSCPPALPAPPRR
		1				CSAGSRTGPAARWCRAGAGSR
						RHAGSSSA***SRR*GRSASGRR
						PRHGGTAGSGRWLWWARCPA
		ŀ		l		*AGIPWPSWPCHSSWP/POSVRV
		1				VSSG*RASLHSPGLDSSGGGLQ
						SFLLVQLPAQGKTSSQP\RHAA
1		1				APSAAACPHAE*CTLGSCPRSA
1		1				TLSPTPPRASPPAG\RSAVSPCPP
		1				GHEGVLVISKHHLLLLSRSGRG
1		1				LGGNVVRPCSVQPMDGVGQEL
			ł		1	AVADRDOLPCLVCQDGECPLK
		1		1		ASGRPVLGCP
L	L		l	L	L	ASORI YEUCI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14727	45095	A	14813	921	1446	ACCISHGNLWHHLDQKHGLQS *GEKSRLFSRPNLQVPVHTGLD HRARYLTGPRKQGHLWPTGPS
					1	SVFDNENFCKLSVGM*TGRASA
						RQQIQSPFAESGKAFSDICSVSL
						RLCVLPHPWNSRLRSPALRNSL
						KYSVTSSAAARRKLLPLASGPL
						RLGPRSLPTRLQVLRWEKVASA
14728	45096	Α	14814	1	675	
14729	45097	A	14815	1	2341	MASDLVNRCEACDLPQGPISVH
						QFPRLSNEVTTTVGRNSESPNQ
1					1	RSVSGPWTPREVRRGVAAAAH
ļ						GFSGQNAKWAAHIPATAMSFA
						LPYSVLHSPHALTTHPLNPGDA
						VTMPSWTSTVSAANASKGRSFS
					1	QCFQSTEEGOERRRPRTSOAKV
						QQGSTAGKVNELITLAGHIKSQ
İ		1			1	APHGKRKPPPRDSHVICSMLRK
						RKPDETPSPSSDAQFSGMLAVA
		l				VSSLLCPRSSPSQQWHCHSPSTT
]	1	ĺ				ILGSSDVSDTSFQPSTKLGPVHG
	l					FPPQSRDCILDFELIRKVTGAWS
						SLTAGSWETKIQEFSEEDILAPH
	1					RITPGPHSSKSPVHFLASLWGR
						KTKLYHHTIQDKEPRSVPEAGS
			l			SGRRDGGGAAHSMLCMAPEPD
						HPGSNPNCRSPATLHADADGN
						LOREAAKCQGDGTQVSTHVDH
						TSAPROAETGPATGENEKMESH
						GRSKRGLVGADQCPAETLDSY
1	İ					TLLVGHKKRDKDRQTPGPAPQ
İ		l		ŀ		VNSFVPENYLLRGPLAPLGSAIS
				}		MKTAGAVHIRPSGDLGRAPAP
						AVRPRPRPPRRPRRLLPGPAHLP
1						GAPRPPAPAPLSRRAARQSPAA
		l				AAAPPPGCE/RRG/PAGPASGAG
						PPAPAHGARPGGKPAANPPRAP
						OP*P/RRRRSRRRAQPRPGPAPL
		1				SRMRAEGPARALRPPPHYOPEL
		1		l		GPPSETVGPVTELLAPIPTARFP
						TRSHPYPVTASPARSS*TSGLD*
14730	45098	A	14816	44	455	HLRNRTR/PSOITPHIYNHLIFDK
					1	/PLFNIWWWENWLAICRKLKLD
l	1					PFLTPYTKINSRWIKD*NIRPKSI
						KNLEENLGNTIQHLG/IGKDFM
l						TKTPKAMATKAKIDIWDLIKLK
l						SFCTAKETIIRVNRLPT*WEKIF
l			1			AIYP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide locatiun of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon,/-possible nucleotide deletion, \-possible nucleotide insertion)
14731	45099	A	14817	1	279	IQDIGMGKDFMTKTPKAMATK AFIDKWDLIQLKSFCTLKEITIR VNRQTTEN\YPSDKGLISRMLP NF*RLFPAS*KKPPLFDIGKTEK TNYS
14732	45100	В	14818	i i	2025	
14733	45101	В	14819	1	3570	
14734	45102	В	14820	1	1380	
14735	45103	Α	14821		2742	KKRESSLTHYMRPASFYQSOA FTOQKKRILOPYP**TLMQKSSI KYYLAKRIQQHIKKLIHHDQVGF IPGMGGWFNIRKSINVIQHINKA KDKNHMISIDAEAFDKIQNG MLATLINKLELEETTYKFIWNQ KRAHIAKSILSQKNKAGGITLR DFKLYYKATVTKTAWYWYON RDIDQWNRTEPSEITLHIYNYLI FDKPEKNKQWGTDSLFNKW WENWLAICRKLKLDFFLTPYTK INSRWIKDLNVRPKTIKTLEENL GITQDIGMGKDFPMKTPKAMA TKAKIDK WDLIKLKSFCTAKET TIRVNRQPTKWEKIFTTYSSDK GLISRIYNELKQIYKKKTNNPIK KWAKDIMKHFSEEDIYAAKKH MKKCSSSLAIREMQIKTTMRYH LTPVRMAIIKKSGNRHAPFSIH THIWFIGLYLIQIQKDLSILGFW YPRGILEPIY YPRGILEPIY YPRGILEPIY YPRGILEPIY

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SEQID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14736	45104	A	14822		1698	MEDEMNEMKQEGKFREKRIKR NEQSLQEIWDYVERPALLIGV PSEDGENGTKLENTLQDIQENF PNLARQTNIQIQEIQRMPQRYSS RRATFRHIIVRFTKVEMKEKMS RAAREKEIGTTIREYYKHLYAN KLENLEEMDKFLDTCTLPRLND EEVESLNRIVTGSEIVAIINSLPT KKSPOPDOFTAEFYQRQSESQI MSEFPFTIASKRIKYLGIQLTRU KVDLFKENFYPLLNEIKEDTKK WKNIPCSWVGRINIVKMALLPK VYTFNAIPIKLPMTIFFTELEKTT FKFIWNQKRAHVAKSILSQKNK AGGITLPDFKLYYKATVTKTA WYWYQNRDIDQWNTEFSEIM LHIYSYLIFDKPEKNKQWGKDS LENKWCWENWLAICRKLKLDP FLTPYTKINSWWIKDLNIRPKTIE TLEENLGTIQDIGMGKDFMSK TSKAMSTKAKIDK WDLIKLKSF CTAKETTIKNNQPILKWEIFA TYSSDKGLISRIYMELKQIYKKK CTAKETTIKNDRYKWENDNNIRYKEEL
						YAAKRHMKKCSSSLAIREMQIK TTMRYHLTPV
14737	45105	В	14823	91	1596	TIMETHEILA
14737	45105	A	14824	1	2757	
14/38	45100	Λ.	14024	l'	2131	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14739	45107	Α	14825	ī	2677	MGKKQSGKTGNSKKQSASPPP
						KEHSSSPAMEQSWMENDFDEI
						NKIDRLLARLIKKKGEKNQIDT
						TKNDIGDITTNPTEIQTTIREYY
						KHLYTNKLENLEEMDKFLDTY
		ŀ	1		1	TLPTINQEEIESLNRTITGSEILAI
		1				INSLPTKKSPGPDGFTAEFYQRY
		1	İ			KEELVPFLLKLFQSIEKEGILLN
		1	l			SFYEASIILIPKPGRDTTKKENFR
			l			PISLMNIDAKIFNKILANRIQQHI
		i	İ			KKLIHHDQVGFIPGMQGWFNIR
		1				KSINVIQHVNRTKDKNRMIISID
		1				AEKAFDKIQQPFMLKTLNKLGI
						DGTYLKIIRAIYDKHTANIILNG
						QKLEAFPLKTGTRQGCPLSPLL
						FNIVLEVLARAIRQEKEIKGIQL
						GKEEVKLSLFADDMIIYLENPIV
						SAQNLLKLISNFSKVSRYKISVQ
		l				IPQAFLYTNNRQTESQIMSELPF
						TIASKRIKYLGIQLTRDVKDLFK
1						ENYKPLLNEIKEDTKKWKNIPC
						SWAGRINIVKMAILPKNWKKT
						TLKFIWNQKRARIAKSILSQKN
						KAGGITLPDFKLYYKATATKTA
				ŀ		WYWYQNRDLDQWNRTEPSEIT
						PHIYNYLIFDKPDKNKQWGKDS
				l		LFNKWCWENWLAICRKLKLDP
i		ļ		l		FLTPYTKINSRWIKDLNIRPKTI
				İ		KTLEENLGITIQDIGMGKDFMS
				l		KTPKAMATKAKIDKWDLIKQE
		l		İ		SFCTAKETTIRVNRQPTKWEKIF
		l				ATYSSDKGLISRIYSELKQIYKK
14740	45108	В	14826	I	1186	
14741	45109	Α	14827	1	2304	
14742	45110	В	14828	1	2995	
14743	45111	Α	14829	I	3095	
14744	45112	A	14830	1	3654	
I4745	45113	Α	14831	1	3189	
I4746	45114	Α	14832	I	2064	
14747	45115	В	14833	II	3031	
14748	45116	Α	14834	I	3192	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
1				sequence		
		Ļ.				
14749	45117	Α	14835	1	3210	MVKGSIQQEELTILNIYAPNTG
1						ALRFIKQVLRDLQRDLDSHTIIM
		1		ļ	·	GDFHTPLSTLDRSTRQKVNKDI
		l		l		QELNSALHQEDLIDIYRTLHPKS
		1				TEYTFFSAPHHTYSKIDHIVGSK
				l		ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
1			i			LNDYWVHNEMKAEIKMFFETN
		ŀ				ENKDTTYQNLWDTFKAVCRGK
					ł	FIALNAHKRKQERSKIDTLTSQL
- 11						KELEKQEQTHSKASRRQEITKIR
		1				AELKEIETO
14750	45118	A	14836	1	3144	
14751	45119	Ā	14837	113	1336	PTLAGTGNQETSHLGFNAIPIKL
14,51	1.5117	ľ`		[]	1	RMTFFTEL/EKTTLKFIWNOKR
		1				ARITKSILSQKNKAGGITLPDFK
		1	}			LYYKAIVTKTVWYWYQNRDID
		1				OWNRTEPSEIMPHIYNYLIFDKP
		1				DKNKKWGKDSLFNKWCWEN
		1				WLAIYRKLKLDPFLTSYTKINS
		1				RWIKDLNVRPKTIKTLEENLGN
		l				
		ŀ				TIQDIGMGKDFMSKTPKAMAT
		ŀ		i		KDKIDKWDLIELKSFCTAKETTI
						RVNRQPTEWEKIFATYSSDKGL
						ISRIYNELKQIYKKKTNNPIKKW
						VKDMNRHFSKEDIYAAKRHMK
						KRSSSLAIREMQIKATMRYHLT
		1		Ì		PVRMAIIKKSGNSSRSNIVHSVT
		1			İ	QANALESSWIPLFLIPHIQYISKP
		1				QASSFKMYHVTDPILPPSSQHL
		1		1		DQAAIINLCLQWPLLWFECMCP
		_				SKIHMLKLTPQ
14752	45120	В	14838	1	3127	
14753	45121	Α	14839	1	2742	
14754	45122	Α	14840	1	1998	
14755	45123	Α	14841	1	3030	
14756	45124	В	14842	1	2558	
14757	45125	A	14843	1	3828	
14758	45126	Α	14844	1	4134	MRKKQSRKTGNSKKQSTSPPPK
	1	1				ERSSSPAMEQSWTENDFDELRE
1		1				EGFRRSNYSKLQEEIETKGQEV
	1	1		l		ENLEKNLDKCITRITNIEKCLKE
	1	1				LMELKAKARELHEECRSLRSRC
		1				DQLEERVSVMEDEMNEMKQE
	1	1	1			GKFREKRIKRNEQSLQEIWDYV
		1				KRPNLPPIDVPESDRENGTKLE
		1		1		NTLQDVIQENFPNLARQANIQI
	1	1		I		QEIQRMPQRYSSRRATPRHIIVR
	1	1	l	I		FTKVEMKEKMLRAAREKAFKQ
		1				ASRREDDIAKVTSG
14750	45127	В	14845	1	3555	NOUNT TOU
14759	45127	D	14043	<u> </u>	3333	<u> </u>

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	
	sequence		09/540,217	codon for peptide sequence	or pepude sequence	deletion, \=possible nucleotide insertion)
14760	45128	Α	14846	1	3345	
14761	45129	Α	14847	I	3451	HTKNPSVHHHRQRPKVDKTTK
						MGKKQNRKTGKSKTQSAFPPP
1	1					KERSSSPAMEQSWMENDFDEL
						REEGFRRSNYSELREDIQTKGK
1				1		EVENFEKNLEECITRITNTEKCL
						KELMELKTKARELREECRSLRN
						WIKSQDPSLCCIQETHLMCRDT
i				1		HRLKIKGRRKIYQANGKQKKA
				1		GVAILVSDKTDFKPTKIKRDKE
						GHYIMVKGSIQQEELTILKIYAP
	1					NTGSPRFMKQVLSDLQRDLDS
	1			Į.		HTLIMGDFNTPLSTLD
14762	45130	Α	14848	1	2606	MVASRSSRVGVFYGGQNSVDG
	1					VSRRSDGSLNELCLYRPPTGAY
				İ		LAYDTLDVLPSGQFTVNTPPID
I				i		VNGKSLLALLWEHTSLLTSMG
	1	1				GGRWAMQPLGQLGQLEHLGD
1	İ	1				RRDQVLDRSGRSRRLDAAQHL
						RHRPDRRPQGMVGVLTNQKEP
	1					RTRWIHSRILPEVQGGTGTVPSE
						TIPIDRKEGILPNLFDEASIILIPK
	i	1				RGRDTTKKENFRPISLMNIDAKI
1	1					LNKILANRNQQHIKKLIHHDQV
1						GFIPGMQGWFNICKSINVIQHIN
1		1				RTKDKNHMIISIDAEKAFDKIQ
						QPFMLKTLNKLGIDGTYLKIIRA
	1					IYDKPTANIILNGQKLEAFPLKT
	1	1				GTRQGCPLSPLLFNIVLEVLAR
						AIRQEKEIKGIQLGKEEVKLSLF
1	1					ADDMIVYLENPIVSAQNLLKLIS
1						NFSKVSAYKINVQKSQAFLYTN
1					İ	NRQTESQIMSVFPFTIASKRIKY
1						LGIQLARNAKDLFKENYKPLLN
1	1					EIKEDTKKWKNIPCSWVGRINI
1						VKMAILPKVIYRFNAIPIKLPMT
						FFTELEKTTLKFIWNQKRARIA
1						KSILSQKNKAGGITLPDFKLHY
	Ì		1			KITVTKTAWYWYQNRDIDQW
		1	İ	I		NRTEPSEITPHIYNYLIFDKPEK
	1	1	1	1	l	NKQWGKDSLFNKWCWENWLA
		1				ICRKLKLDPFLTPYTKIHPRWIK
	1	1	1	I		DLNVRPKTIKTLEENLGNTIQDI
	1	1	1	1		GMGKDFMSKTPKAMAAKAKI
		1	l			DKWDLIQLKSFCTAKETTIRVN
14763	45131	Α	14849	1	2862	
	1	11.			l	L

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleoride
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14764	45132	Α	14850	I	3724	MGKKQNRKTGNSKTQSASPPP
			l			KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELWEDIOTKGKE
						VENFEKNLEECITRITNTEKCLK
						ELMELKTKARELHEECRSLRSR
1						CDQLEERVSAMEDEMNEMKRE
			ĺ			GKFREKRIKRNEOSLOEIWDYV
1						KRPNLRLIGVPESDVENGTKLE
						NTLQDIIQENFPNLARQANVQI
			1	1		QEIQRMPQRYSSRRATPRHIIVR
			l			FTKVEMKEKMLRAAROKAPH
						HTYSKIDHIVGSKAL
14765	45133	Α	14851	2	2375	
14766	45134	A	14852	1	2376	MGDFNTPLSTLDRSTROKVNK
11100		,		1		DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKGREIITNCLSDHS
			ŀ			AIKLELRIKNLTQNRSTTWKLS
						NLLLNDYWVHNEMKAEIKMFF
1	ļ		i			ETNENKDTTYONLWDTFKAVC
1				ļ		RGKFIALNAHKRKQERSKIDTL
1			İ	l		TSOLKELEKQEOTHSKASRROE
				1		ITKIRAELKEIETOKTLOKINESR
				İ		SWFFERINKIDRPLTRLIKKKRE
			İ	1		TNOIDAIKNDKGDITADPTDRO
ŀ						TTIREYYKHLYANKLENLEEM
ļ						DIFLDTYTLPRLNQEEFESLNRP
1				i		ITRSEIVAIINSLPTKKSPGPDGF
				1		TAEFYQRYKEELRIKYLGIQLT
1				1		RDVKDLFKENYKPLLKEIKEDT
				ŀ		NKWKNIPCSWVGRINIVKMAIL
				1		PKVIYRFNAIPIKLPMTFFTELE
			İ	1		KTTLKFIWNQKRARIAKSILSQ
				1		KNKAGGITLPDFKLYYKATVT
			ŀ	1		KTAWYWYONRDIDOWYRTEP
			1	l		LEITPHIYNYLIFDKPEKNKQW
			1	l		GKDSLFNLWCWENWLAICRKL
1	l			l		KLDPFLTPYTKINSRWIKDLNV
1				l		RPKTIKTLEENLGITIQDIGMGK
1	l		1			DFMSKTPKAMATKDKIDKWDL
1	l		1			IKLKSFCTAKETTIRVNROPTK
1			l			WEKIFTTYSSDKGLISRIYNELK
1				1		OIYKKKTNNPIKKWAKDMNRH
			1	ŀ		FSKEDIYAAKKHMKKCSSSLAI
						REMOIKTTMRYHLTPVRMAIIK
L	Щ	\Box	Ь	L		Little Control of the

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14767	45135	A	14853	I	1576	MDKFLDTYTLPRLNQEEVESLN
		1				RPITGAEIVAIINSLPTKKSPGPD
					\	GFTAEFYQRLISNFSKVSGYRIN
						VQES\QAFLYTINRQTESQIMSA
						LPLTIASKRIKYLGIQLTRDVKD
						LFKENYKPLLNEIKEDTNKWK
i						NIPCSWVGRINIVKMAILP/KEL
						EETTLKFIWNQKRARIAKSILSQ
						KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDRDQWNRTEP
						SEIMPHIYNYLIFDKPEKNKQW
						GKDSLFNK WCWENWLAICRKL
	1	1				KLDPFLTPYTRINSRWIKDLHV
		1				RRKTIKTLEENLGNTIQDIGMG
		1				KDFMSKTPKAMATKAKIDKW
		1				DLIKLKSFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYNEL
					1	KQIYKKKTNNPIKKWAKDMNR
						HFSKEDIYAAKKHMKKCSSSLA
						IREMQIKTTMRYHLTPVRMAII
						KKSGNN/R/CAPGTPERQNHSLS
						WKGS*SQEPSGLAQWIPLPWSP
	1	i				AS*DPLA*NSCCQHSCLKSTWD
						AQACARKLRTLIKGYRNCQLE
14768	45136	В	14854	328	1467	
14769	45137	Α	14855	3	3412	
14770	45138	A	14856	1	2907	MDKFLNTNTLPRLNQEEVDSL
		1			\	NRPITGSEIVAIVNSLPTKKSPGP
1		1				DGFTAEFYQRYKEELRIKYLGI
ĺ		1				QLTKDVKDLFKENYKPLLKEIK
		1				EDKNKWKNIPCSWVGRINIMK
		1				MAILPKVIYRFNAIPIKLPMTFFI
		1				ELEKTTLKFIWNQKRALIAKSIL
		1		1	1	SQMNRAGGIMLPDFKLYYMAT
	1	1		1	1	VTKTAWYWYQNRDIDQWNRT
	1	1		1	1	EPSEITPHIYNYLIFDKPEKNKK
		1		1	1	WGKDSLFNKWCWENWLAICR
l	1	1	1			KLKLDPFLTPYTKI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14771	45139	A	14857	1402	3004	TEPKIKTI*LSA*MOKGPLTKF NNPSC*KLLINIVLEVLARAIRQ EKEIKGIQVGKEEVKLSLFADD MIVYLENPIVSAQNILKLISNIS* CVSGYKINVOKSQAFLYTNIN QTESQIMSELPFTIASKRIKYLGI QLTRDVKDLFKENYKPLLNEIK EDTINKWKNILCSWVGRINIVK MALIPKVIYRFNAIPIKLPMPFT ELEETTLKFIWNQKRARIAKAII. SQKKKAGGTILPDFKLYYKAI'V TKTAWYWYQNTDIEOWNRTE SEIMLHIYN/LIFDQPEKNKQW GKDSLFNKWCWENWLAICRKL KLDPPLTPYTKINSRWIKDLNV RPKITKTLEENIGITIQDISNOK DFMSKTPKAKATKAKAIDKWDL IKLKSPCTAKETTIRVNRQPTK WEKIFATYSSDKGLISRIYNELK QIYKIKTINPIEKWVKDMNRHI SKEDIYAAKKHMKKCSSSLAIR EMQIKTTMRYHLTAVRMAIIKK SGNNSLIVGASMDTSLAGIPFY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14772	45140	A	14858	884	2925	GIKYLGIQLTRDVKDLFKERS/Y
				ŀ		EPLLNEIKEDTNKWKNIPCSWV
l						GRINIVKMAILPKVIYRFNAIPIK
1		l				LPMTFFTELEKTTLKFIWNQKR
		l				ALIAKSILSOKNKAGGITLPDFK
		1				LYYKATVTKTAWYWYONRDI
l		1				DOWNRTEPSEITLHIYNYLIFDK
						PEKNKOWGKDSLFNKWCWEN
		1				WLAICRKLKLDPFLTPYTKINSR
		l	İ	ŀ		WIKDLNVRPKTIKTLEENLGITI
		Į.		ŀ		ODIGMGKDYMSKTPKAMATK
	ŀ					AKIDKWDLIKLKSFCTAKETTIR
	i i					VNRQPTKWEKIFATYSSDKGLI
				l		SRIYNELKQIYKKKTNNPIKKW
	li .			l		VKDMNRHFSKEDIYAAKKHM
	1			l	1	KKCSPSLAIREMQIKTTMRYHL
				l		TPVRMAIIKKSGNNSPTCVSQD
				i		WKLTQAVGLWSAEAEAPKSAQ
	1	l				RRPLAEGHRAAVQRDGSGLQL
	i					LCTVKLQVLPCIIAQGNGGRKA
	1	ĺ				TSSLEAELTSWRPLCLEDGTGF
	ŀ					RGEDTLETVNAEGRENRRKAG
l	i			i		HRAAGSSASRRCRNTGRSVQRL
1	1					RPQKTQAFPTHCHFCLILVVKA
	1					TQLNPKSRVRKVPFAPMKHGK
l	1			1		ALAGPWANICAGKSSNEIRTCD
						RHGCGQYSAQRSQRPHQGVDI
	1					LCSAGSTVYAPFTGMIVGQEKP
						YQNKNAINNGVRISGRGFCVK
l	1	1		i		MFYIKPIKYKGPIKKGEKLGTLL
						PLQKVYPGIQSHVHIENCDSSDP
14773	45141	Α	14859	1	2577	

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SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1	1	l		sequence	' ' '	
14774	45142	Α	14860	3019	5574	RETRGEVKNSKNQSTSAPPKDC
		l				SSLADPKKIQTTIREYYKHLYPN
	l	1				KLENQEEMDKFLDTYTLPRLN
				i		QEEVESLNRPITGSEIVAIINSLP
						TKKSPGPDGFTAEFYORYKEEL
		l				VPFLLKLFQSIEKEGILPNSFYE
		1				ASIILIPKLGRDTTKKENFRPISV
		1				MNIDVKILNKILANRIOOHIKKI
		l				IHHDOVGFIPGMQGWFNIHKSI
		l				NVIQHINRTKDKNHMIISIHAEK
						AFDKIQOHFMLKTLNKLGIDGT
		l				YLKIIRAIYDKPTANIILNGQKL
		1				EAFPLKIGTRQGCPLSPLLFNIV
		1				LEVLARAIRQEKEIKGIQLGKEV
		1				
		l				VKLSLFADDMIVYLENPIVSAQ
		1				NLLKLISNFGKVSGYKINVQKS
		l				QAFLYTNNRQTESQIMSELPFTI
		l				ASKRIKYLGIQLTRDVKDLFKE
	1	1				NYKPLLNEIKEDTNKWKNIPCS
		1				WVGRINIVKMAVLPKLPMTFFT
		1				ELEK\TTLKFIWNQKRARIAKSI
		l				LSQKNKAGGITLPDFKLYYKAT
		1				ATKTAWYWYQNRDLDQWNRT
1		1				EPSEITPHIYNYLIFDKPDKNKQ
	İ	l				WGKDSLFNKWCWENWLAICR
		l				KLKLDPFLTPYTKINSRWIKDL
		1				NIRPKTIKTLEENLGITIQDIGMO
1		I				KDFMSKTPKAMATKAKIDKW
		1	l			DLIKQESFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYSEL
1						KQIYKKKTNNPIKKWAKDMNR
		l				HFSKEDIYAAKKHMKKCPSSLA
14775	45143	Α	14861	1895	6140	
14776	45144	Ā	14862	I	2875	MKAEIKVFFETNENKDTTYQNL
1	1	1		l		WDTFKAVCRGKFIALNAHKRK
		1				QERSKIDTLTSQLKELEKQEQT
		l				HSKASRROEITKIRAELKEIOTO
		1		l		KTLOKINESRSWFFERINKIDRS
		l		l		LARLIKKKREKNQIDTIKNDKG
						DITTDPTEIQTTIREYYKHLYAN
		1		1		KLENLEEMDKFLDTYTLPRLNO
		l				
		l	1			EEVESLNRPITGAEIVAIINSLPT
		1				KKSPGPDGFTAEFYQSWAETQP
1	l					KKENFRPISLMNIDAKILNKILA
L	L	L	l	L		KRIQQHIK

SEO ID	SEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
14777	45145	Ā	14863	2843	4460	HQHQQKPHPHRNPIQKSPTSKT
14	13.13	1.	1.1005	20.5	1	KVLEVLARAIRQEKQS/GIQIGS
						QEVKLSLLEDDMIVHLENPIISA
Į.						KNLLQLISNFTKVSGYKINVQK
						SQAFPYTNKTQIENQIMSELPFT
						IATKRIKYLGIKLTRDVKDLFK INNYKPLFKEIREDTNKWKNIPC
		1				SWVGRINIVKMAILPKVIYRFN
		Į.				AIPIKLPMTFFTELEETTLKFIW
1						NQKRARIAKSILSQKNKAGGIT
						LPDFKLYYKATVTKTAWYWY
						QNRDRDQWNRTEPSEIMPHIYN
						YLIFDKPEKNKQWGKDS\LFNK
1						WC*ENWLTICKRLKLDPFLTPY
		1		1		TKINSRWIKDLNVRPKTIKTLEE
1		1		1		NLGITIQDIGIGKDFMCKTPKA
		1				MATKAKIDKWDLIKLKSFCTA
		1		1		KETTIRVNRQPTKWEKIFATYS
1		1				SDKGLISRIYNELKQIYKKKTN
1		1				NPIKKWAKDMNRHFSKEDIYA
1		1				AKKHMKKCSSSLAIREMQIKTT
		1				MRYHLTPVRMAIIKKSGNNRC
						WRRCGEIGTLLHCWWDCKPVQ
			İ			PLWKSVWRFLRDLELEIPFDPAI
14778	45146	A	14864	I	8335	PLLGIYPKGYKSC MGKKQNRKTGNSKMQSASPPP
14//8	45146	A	14864	1	8333	
1						KERSSSPATDQSWMENDFDEL REEGFRRSNYSELREDIQTKGK
1	1		l			EVENFEKNLEECITRITNTEKCL
						KELMELKTKARELREECRSLRS
						RCDQLEERVSAMEDEMNEMK
i	1		i			OEGKIKRDKEGHYIMVKGSIOC
			1			EELTILNIYAPNTGAPRFIKQVL SDLQRDLDSHTLIMGDFNTSLL
	1					
				l		TLDRSMRQKVNKDTQELNSSL
	1					HQADLIDIYRTLHPKSIEYTFFS
14770	45147	 -	14865	1	3166	APHHTYSKIDHIVGS
14779	4314/	A	14860]'	13100	MVKGSIQQEELTILNIYAPNTG
		1		1		APRFIKQVLSDLQRDLDSHTLI
1						MGDFNTPLSTLDRSMRQKVNK
				1		DTQELNSALHQADLIDISRTLHF
		1		1	1	KSTEYTFFSAPHHTYSKIDHIVG
1		1		1	I	SKALLSKCKRTEITNYLSDHSA
1		1		1	1	IKLELRIKNLTQSRSTTWKLNN
		1		1	I	LLLNDYWVHNEMKAEIKMFFE
				l	l	TNENKDTTYQNLWDAFKAVCR
		1		1	1	EKFIALNAYKRKQERSKIDTLTS
	1	1			1	QLKELEKQEQTHSKASRRQEIT
	L	1				KIRAELKEIETQ
14780	45148	A	14866	I	3912	
14781	45149	В	14867	I	2832	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or pepuae sequence	deletion, (-possible nucleotide insertion)
				sequence		
14782	45150	Α	14868	1944	3531	LRDCKRINQHPVKTDQSAFCK
						M/API/MQDVVLEVLARAIRQEK
						EIKGIQLGKEEVKLSLFADDMI
						VYLENPIVSAQNLLKLISNFSKV
						SGYKINAQKSQAFLYTNNRQTE
			1			SQIMSELPFTIASKRIKYLGIQLT
		1		į		RDVKDLFKENYKPLLKEITEDT
		1				NKWKNIPCSWVGRINIVKMAIL
						PKVIYRFNAIPIKLPMTFFTELE
		l				KTTLKFIWNQKRARMAKSILSQ
		1				KNKAGGITLPDFKLYYKATVT
		l				KTAWYWYQNRDVDQWNRTEP
		1				SEITPHIYNYLIFDKPDKNKQW
						GKDSLFNKWGWENWLAICRKL
		1		i		KLDPFLTPYTRINSRWIKDLNV
1		1				RPKTIKTLEENLDITIQDIGMGK
1						DFMSKTPKAMATKAKIDKWDL
i i						IKLKSFCIAKETTIRVNRQPTKW
i						EKIFATYSPDKGLISRIYNELKQI
				i		YKKKTNNPIKKWAKDMNRHFS
1				1		KEDIYAAKKHMKRCSSSLAIRE
		l				IQIKTTMRYHLTPVRMAIIKKSG
		l				NN/R/CF*WQKPGPSG*ALRSWK
				1		EDIAGTSYVGMNQCQMRKNTK
						TLI
14783	45151	Α	14869	1182	3891	KMIKGISTTDLTEIQTTIREYYK
i					\	HLYANKLENLEEMDKFLDTYT
ĺ		i				LPRLNQEEVESLNRPITGAEIVA
						IINSLPTKRSAGPDGFTAEFYQR
		1				YKEELVPFLLKLFQSIEKEGILP
						NSFCEASIILIPKPGRDTTKKENF
i						RPISLMNIDAKILNKILAKRIQQ
						HIKKLIHHDQVGFIPGMQGWFN
						IRKSINVIQHINRAKDKNHMIISI
		1				DAEKAFDKIQQHFMLKTLNRL
ļ		1				GIDGTYFKIIRAIYDKPTANIILN
						GQKL
14784	45152	Α	14870	997	1161	CTSTVIMSICTKLQNKEHVIEAL
1		1				RRAKFKFPGHPK\IHISKK*DFT
1		1	[KFNADEFED
14785	45153	Α	14871	I	1458	
14786	45154	В	14872	53	1468	
14787	45155	Α	14873	1819	2235	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14788	45156	Α	14874	707	1613	VRARAGERARVVRENGKAELP
						VALVLPSRALSHTFTCTDLKVQ
						FHQRLRLQEKGSKFIGSNMSQE
						SDNNKRLVALVPMPSDPPFNTR
				l		RAYTSEDEAWKSYLENPLTAA
						TKAMMSINGDEDSAAA/HRPAL
						*LLQENQSEPIPREGVGWG*GE
						ER\ERMTEEEQEGEGGGGRGG
						GEGEEEEEEEDVEEEEEEEEE
						EKEEEEKGGEKEEENEEKEKQR
						EGRRGKETSNGQMVRPRLLPP
						HQEHSRNYLVFIMDSNMIEICF
						REASISPKAQAYVLIPDFKNCIY
					1	GWEEDSIEILWGLPGSHVL
14789	45157	Α	14875	3	346	QRSKSCKVPGLGKPLTLPPKPE
						K\PQGQKDRRPTGFKP*N*RRRR
						SEKSLRFFPPGS/PQAVPIPVGFS
			1			FDTNL*HFL*KPLIFIMGGTVSL
						PHPRSPSAPPGTPQRPMTFCHLK
		l				Q
14790	45158	Α	14876	1868	2666	GLLLSPFQGLQGVAPRKTRSNL
1						HPGSLAHLSHQPPAPWPPRGLW
		l				PPGQETSCHCASGRALGTGSPR
						PPHAPWLHLEPQRAPHSGAPSP
l					ì	GSSPSAPLHIPVAPENLQSLGHR
						SPPNPSGPDCYPSPSGP/PPSSLD
						SPGLQTAPGAMFLSPRSGVQSA
						CPTPRSLDLRNSPASTPDQSVCP
						TPLSLAWLWPPPSPTSAKSLLFT
						LTSEVQSFSPTPIPPGSSDPPLSK
						QP\SGSCLSSSGHLTSPSVPRHSP
		ļ. —				WPRSSPQNPHSWTLPSS
14791	45159	Α	14877	762	1528	AIPGGAAWPFRAEGAQGPGKR
						LTFKPLGAQRQSVSGGSEVPPL
						PLHLKMAPEMPVAL*SPGTGV
						HPSLLPPT/GPKVPPGRIDRGPVP
						TLAPQVGTAPIQWHCRGSRCPS
	1			l		G/SRPPGGWR/RPCQAPAPPLGQ
				1		HLPRLLPPPTGPQGPRPLTPSEP
1				1		PPWSCWLPASPPAGPGSRRVGG
		1	l	l		GPGTCQGLLRS*GLLLSPFQGL
				1		QGVAPRKTRSNLHPGSLAHLSH
				l		QPPAPWPPRGLWPPGQETSCHC
L	L	L		L	L	ASGRAVGTGSRRPRA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	Incation of first	codon for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequence		
14792	45160	A	14878	1	963	MARKDCCSPPGDSSPGRLPLVI.
14772	45100	l^	14070	Ι'	J***	PAOCLVLLRLOITAFSSIHVAAN
						DMMSFFLMAKQDSIAYKYRIFF
		l				IHSSVDGHARPLSPSLVVENGC
		l				GKOVFLLCSDNASPEVSTVALA
1						SCPGTFPRHCGAAESTHICATG
		l		1		LGELTPOGHGHIPNKLCACCAT
l .						GLEELVPRDHGKLGISWFGFIA
1		1				GAYLNLARVVGECOGSYGYLS
		ŀ		1		LHHYCYCHYYHYCHYHYH\AA
i		1				TY*NHYYHYYH/HHNNSYHHH
						SYYHYSYHFYHYYYNCYHHH
						YHCRHYYHPHDYYHYHA/RHY
1	111					SLYYCHYLCHYYYRCYHDHSS
1				1		HQHNHHECLP*VRHSAEHLQY
14793	45161	A	14879	105	1448	CHOPPASPFSRREPGGSEKPPER
						GCPAWEPAAOGPVOPLGEDGP
						RLCOPPPPSPDHALG\PYGHORS
						GRPTLPGPRK\PAKKRGR*PSSS
						TGQ*SHRPTTGTPGGQAGGSG*
						VTAGGQ*PPQNL*AKISCPEMP
		1				GPSPGMQQGVKHPSLFSHQTV
	l					AKGFTTRWNCRQVCPHRGERK
						WISSVRPCGGALAPSCA/DGLR
		1				GRRTLSGSQPLW/PPAAVSAVG
						QLQPPCHTLTPHPLQNSTQHAE
						PQRGSVYSL*EGTSP\LAAVPFA
						PHIEKRPFSTFPAGDA*PSEAHA/
						PGPACCHSPGGLEGPVSRAGPG
						TS AGR\PGNGRERA WPPHSGPS
	l					TPPDTSCG/SPGPQLARAGSRRA
						NGSTLPPTRQGFLPGRKDSAGA
				1		CGQP*AEGAAAGPSRSPAAER
			1		1	AVLPLPR\GRRKCRGGGPGVET
			1	l		GVSGARRGGDRSGCSPAALDA
		L				EAPASLAGVSIREG
14794	45162	Α	14880	1	2421	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14795	45163	A	14881	1080	2571	VHCRFWILALCQMSRLQKSPLL
						FNIVLEVLAKAIKQEKEIKGIQL
	1					GKEEVKLSLFADDMIVYLENPI
			İ	1		V/SAONLLNLISNFSRVSGYKIN
				1		VOKSOAFLYTNNROTESOIMSE
						LPFTIASKRIKYLGIOLTROVKD
						LFKENYKPLLNEIKEDTKKWK
i				1		NIPCSWVGGRINIMKMAILPKVI
				ļ	i .	YRFNAIPIKLPLTFFTDLERTTL
	l					NFICNOKRAPTLPOVOSLS\OKN
				Į.		KAG\GIT\LPDFKLYLQGLQ*PK
l				1	1	TAWYWY\ONRDIDOWNRTEPS
1				l		EIMPHIYNYLIFDKPWEK/HKO
i		1		İ		WGKDSLFNKWCWE\NWLAICR
				l	ļ	KLKLDPFLTPYTKINSRWINDL
	1					NVR\PKTIK\TLEENLGIT\IOD\IG
			i			MGKDFMSNKTPKA\MA\TKAKI
1			Ì		1	DKWDLI\KLKSF\CTAKETTIRV
						NR\QPTKWKKIFTTYSSDKGLIS
	i					RIYNELKOIYKKKTNNPIKKWA
		l	i	l		KDMNRHFSKEDIYAAKKHMK
	i					KCST\SLAIREMQIKTTMRYHLT
ł		ı				PVRMAIIKKSENSRCWR
14796	45164	Α	14882	136	420	RESRNLSRGEESEDPAEPS\RNG
		1				*EHPLCYLQ*EFLQVLTMLQAE
		l			1	GTMHHFRSICQVNRNFLERGH/
		Ì				SPPSPAPPPETHTGSPRPPSGRSR
						IRAYLH
14797	45165	В	14883	I	1197	
14798	45166	Α	14884	3	538	GAGTDGGRTPAPRAFPDPLGEA
	İ				1	RTSAPTGPARPGGSASAARPAA
						VRPAEPA AEET QEQPRAGKRAI
l	1	1		l	1	HEYRYVYVQQRRKTDPQPAVA
ŀ		1		l	1	VFGIPADGD*VSRTAYHAHQPL
1	1			I	1	AWSGR/E*MCLSMAMWNT*ST
1	1	1		l	I	MKK*ISIRVISRCSGPVHRTN*QI
1		1		l	1	PEPVRAWRFLICRCICFSPGRWI
1	1	1		1	1	KT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino aeid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*-Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nueleotide insertion)
14799	45167	Α	14885	1208	2765	KMAPTWDKMVSSAQMGFNLQ
						ALLEQLSQDELSKFKYLITTFSL
						AHELQKIPHKEVDKADGKQLV
						EILTTHCDSYWVEMASLQVFEK
						MHRMDLSERAKDEVREAALKS
		l				FNKRKPLSLGITRKERPPLDVDE
						MLERFKTEAQAFTETKGNVICL
						GKEVFKGKKPDKDNRCRYILK
		1				TKFREMWKSWPGDSKEV/RGY
						G*EIQDADPIQOPQGASRAL\PY
				i		TVVLYGPAGLGKTTLAQKLML
				!		DWAEDNLIHKFKYAFYLSCREL
		Į.				SRLGPCSFAELVFRDWPELQDD
						IPHILAQARKILFVIDGFDELGA
						APGALIEDICGDWEKKKPVPVL
						LGSLLNRVPLCIENKLGVGGED
						WGDQLRVYWWNPSKREGGSG
		1		Ì		WSHGSEDANQSLKHLRLSANV
ł				ł		LLDEGAMLLYKTMTRPKHFLQ
		l				MLSPWETTLAQIIMLDWAEETS
	1			İ		SDKLKYAFYCSCRELSRLCPCN
						FEELVFRDWPELQDDIPHILAQ
				l		ARKILFVIDGFDPLGGRTLWTL
				1		TRGHLRELGNRRKRCPSSWGV
14800	45168	Α	14886	213	625	EPGIDTSCLCAGRCSRPGKGRA
						CGGSSPCSWPGQPPAPPAAAAV
						PACPSGGRGAGPIQSGTCCSRS
1						ADSRCTSWTAW/WPPGLGAAT
		1				VQTVMS*TV/WPMPHVPQAGW
		1		ľ		LALGQGAAAAKHPLTSATTRS
		1				WCLGVGPGPWM
14801	45169	Α	14887	I	375	MCNAQVLEYMGKSSSLLTSDL
		1				QLVRDALRSLRNSFSGHDPQHH
						TIDSLEQGISSLMERLHVMETQ
l		1				KKQERKPPLVIPRQT/WVWSGP
l		1		l		PANSNRPAAEGPDC*KEN*RDA
		1		ĺ		LRSLRNSFSGHDPQHHTIDSLEQ
		1		l		GISSLMERLHVMETQKKQERKP
		1				PLVIPRQTGSGVDLQQTPTDLQ
		1		l		LRDLTVRRKTNEQKGIASASTK
l		1		ŀ		RTSTPKPHL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
l				Sequence		
14802	45170	Α	14888	1	750	MGELNQGSWRQGHLPLDAAN
İ						GIAIRVPRRRKDESRKNPTRKPH
						SWKPTKNFEDDVEVMKRSMSI
	1					RIEDTHSGPSAAGVLEFAGGPL
						QTLFAWVSPAEAAEQQRLLPVP
						/CLWKLCPRGAPARC\RQNSPSK
						LPSGFVYTVRGKLPTQASVVVD
		1		į.		APPATKLECPRSTSDCCAGSKN
	1	l				FKPVDLSLLASVGPRVNLQQES
						SHYGKISVTVAGHCPTHATGRI
						LGKQALRWSPMCSMFIGRALG
		ł		l		LVPVEGGLLQIA
14803	45171	Α	14889	81	164	VGKQGLEWTPANSSRPAAEGP
						DC*KEN
14804	45172	Α	14890	1	674	MVVYACSPSYLEAEVGGSLSLE
	i i	1				VEATNPKNSSKKFLDLISEFCK
				ĺ		VSGYKISVHKSVALLYTKNDQ
						AATQIKNLIPFTTAAKQFKYFGI
						YLTKEEERLKPSRKKITKKHTK
	i	l				KRTASLILHAMICSQIPKQQQNE
				I		KYQVPQFDQSTIKNIESAKGLD
		l		i		VWDSWPLQNADGTPPLVIPRQ
	i	1				T/WVWSGPPANSNRPAAEGPDC
	l l					QKEN*QTERNSINIKKKDIHTKT
						PLCI
14805	45173	Α	14891	619	931	YLLMDFPLLHSWLVSLGDKVP
		İ				HEFWRGCPSCSFGPRFWVAGV
				1		LVL*APKL*LRQVRSAGKRCGQ
		l		l		KW*IK*/SDRLCVD*NK/WSDSD
						LKPATSCSLIVTCHSLLGF
14806	45174	В	14892	1	658	
14807	45175	A	14893	48	288	LCSIKKPEKSLLCW*YPGNRVW
				l		SGPPAKSNRPAAEGPDC*KEN*
		l		İ		QTERNSININKKDIHTKTPSEGH
	L	_				QHQRPKTKPGRI
14808	45176	A	14894	801	1157	QSGPSAAGLLDFAGGPLQTLFA
		l				RVSPAEAAEQQRLLPFPSSGKL
						RPRGAPARCQPELSCMRC\R*SL
						LGGVSQSGYMGVRDPFEEAAC
		1				PLSELECSAGRSAALLRMVRQG
İ	_					HLGLLKLRP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amine acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14809	45177	Α	14895	1	1891	MPVWVTVVKSWNTQIGRPGH
						VSALGFELTKNPWPLTYESGDG
				l .		RSLGKPGAVSRRGGRHGPHPQ
			1			QAECDLPAGRIAGPCERGLERQ
				ł		EKLPPAPQGGAWHLRLSRERFK
						SEDAPKIHVALGGSLFLLNLAF
	1			į.		LVNVGSGSKGSDAACWARGA
						VFHYFLLCAFTWMGLEAFHLY
						LLAVRVFNTYFGHYFLKLSLVG
						WGLPALMVIGTGSANSYGLYTI
				l		RDRENRTSLELCWFREGTTMY
				ł		ALYITVHGYFLITFLFGMVVLA
						LVVWKIFTLSRATAVKERGKN
						RKKVLTLLGLSSLVGVTWGLAI
	ļ.					FTPLGLSTVYIFALFNSLQAHKQ
						LKDRKVGPHFMQGAVFLEGKF
İ	ŀ			i		WGVDIASFEPQEQFWWTNLGK
		ŀ				EDEDKSFDMPHSWVEQIEISPEE
				l		EAMGSAQMVRGVSGPERGSKK
		l		1		RPQLTVSTAFETRCPNGKKVIQ
						YKRAKLEKWAPYLNSNGLVSR
				l		LTTYEDLQCTNILEIKEWYQNR
	1					EDMLELKHINKTTDLKTDYFKP
						GHPQALRVHSYKSMQPEMDRV
l		l				IEFYETARVDGLMKREETPRTM
1				l		TEYYQGRPDFLSYRHASFG/RPS
						QEAHSEQCRVKPPAHCAKPAE
						DDV/RKSACFWSRRSASSCATT
						AVRTTSRPPSASSCGAPRWTAK
14810	45178	A	14897	1	425	HVQQTEISRTVSTGSPIILSCSVP
		ŀ				VTLPNGPVLWFKGN/AGPNRKL
				ŀ		IYNFKQGNFPRVKEIGDTTKPG
						NTDFSTRIREISLADAGTYYCV
				l		KFIKGRAIKEYQSGRGTQVFVT
				ŀ		VSYSSRSQKSEICISGLKSRHQQ
						NCSFWRF
14811	45179	Α	14898	J.	393	RQGNFPRAKQIGDTTKPGNTDF
1		l		[STRIREISLADVGTYYCVKFIKG
1		1		ł		RAIKEYQ\SGRGTQVFVTEQNP
						RPPKNRPAGRAGSRAHHDAHT
1		1	1	l		CLSALPERNSTNYFVQPCCCLR
				L	L	LLGLTGFAVKIIQTGKERTSK

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
					L	L
14812	45180	A	14899	1	753	MGSSLSPVLYVALPVRQPLLRS
						VVEERSTKMRVGSPGFQAKFRF
i		l				YSQGLSWSARGEEGLVPAFWL
					l	QPYPRPLNFKAEMTEQKENQT
			l			YVDGGFRPYIHRRKNLTEGHK
						AEKEMEPWDGFSDESCRTHCQ
	ŀ					VHDYAKGHGEVQRERVDEQK
1			l			EHLGGRRQDCQQQHLNRRLHC
						/TVSGEAPSEASGSVRGHGRAH
						GLAGSERHGHDHAAPEHOLLO
				1		SVDADHVVPVAWYOFWDDEG
	ŀ					RGILSPGDHKPDRRGAALY
14813	45181	A	14900	6	726	VEGISRTGIQAEQAPGFTTMPIP
14813	45181	A	14900	ľ	720	
		1				VSVNPGDVLTLACNMSALSPE
1		l				GPVLWFKSIGPGQQLIFSCNGS
1				ŀ		HFPRVTPVENTMVDQTDYSIILS
		l				CSVPNTLPNGPVLWFKGTGPNR
		l				KLIYNFKQGNFPRVKEIGDTTK
	l			ł		PGNTDFSTRIREISLADAGTYYC
	l	1	1			VKFIKGRAIKEYQSGRGTQVFV
		l		l		TGEYVSYTPYPLVYNS\PLWDE
		1		l		ASRGRSRQQSLLFCSLHW*YPR
						KQDLEWTSSKLQQTCRRVA
14814	45182	Α	1490 I	760	978	KKLLLLLCGIEWHGFSSDSGS
		1	ŀ	l		GGWVPEG*K*GCEHELLPEDVP
		1		l		SCFGWAECWLVRRTLPLFHYV
1		1				ELCYSFHG
14815	45183	В	14902	772	1165	
14816	45184	Α	14903	1	2133	
14817	45185	Α	14904	1	1311	MATTPGRALGSLLPTLLLGLTA
1		l				GOPFISLTGPSHRTSPGDSVPFN
		l				CTAVPFNSQDFSRDFNVTWLK
		l				DSDEHPASAQRLVPDNGGNDFI
		l				TSKAWVTLTRODVSSEITCEVT
		l				HRALAEPLKTTMNLSQATRNP
		l		ŀ		DGTYSLEHTWQAEATLEEREF
		l				ACWVVHDEQPPLKANVTLQAQ
		l	į.			VRRQGKGQSVQPYKLQGPLQR
		l	1			SEPGTRIRLTYASSGFSTHQVTV
		l				
		l		}		TWLKNKHELPNPQTSVQYSGH
		l	1	1		TYNVTSSVLVPLMDDDVFSHV
	l	l		1		VCHVEHKLTWFFQKTTGLDQY
		l		l		L/QG/DP*QLCGGHLTLPSSPRAP
1	l	I	i	l		PQLPGLQHP/VVFHSQVLLFSCL
		1		1		VAVTCLVQRFYPQNVHLAWLE
	l	1		l		DCHTLKGTEQPTLKKNNDRSY
1	l	I		1		TLEKLLLVNASVQGPERVLTC
		l		ŀ		MMEHEGQPPKRANLVLSTAAH
	l	l		1		TAYKPIGSTESIMVTMIITISQM

SEQ ID			SEQ ID NO:			Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or pepade sequence	desertion, v-possible nucleotide insertion)
14818	45186	Α	14905	3	341	RRATRPGN*GGRHNAGRSSARL
						ETGRPAMAPQGRPRPLGCRTPO
						HPGHWPSAPGLLPPAPGPVAG
						MTQAQRWCPEPDAHFLQYTLY
						FVRHIPQLPKQTPHHYIATRLPC
		į				ATQ
14819	45187	Α	14906	154	679	GVCCPLLGGVSS*ATWGSGTT*
						GGNTHLI
14820	45188	В	14907	1	2554	
14821	45189	Α	14908	387	826	ARAACLPRGSARGTAGAGEEPP
1						DPAPLHLRGEGPAP/HRPRSPRG
1		1		1		MGWPAAPSTRSPPST*RRCPLPP
1				1		RPRWSP*SHPGPRASARRLP*LS
						*RASRSSEAPSPGSLGSVAGQD
						CSRTGTGFSPPAFPSDGQQTPPP
				i		APSSVPTLNSLG
14822	45190	Α	14909	86	2223	GQEPQVREQKACHHLGSSPPPS
1			İ			WEL*EQRPAGP*N**PSTLTTGT
						GSTTT*QIRMKVN*MGN/DSPQ
						GKNTPKMYSGEFSPVRVHVPFS
						LSDLKQIKIDLGKFSDDPDGYID
						VLQGEDILAKAGAIRHLNIGEG
1						THVCCPLLEEGINPEVWATEGK
1						YGRAKNAHPIQVKLKDSASFPY
1						QRQYPLRPEAQQGLQKIVKDL
1						KAQGLVKPCSSPCNTPILGVQK
			1	l		PNGQWRLVQDLRIINEAIVPLY
1						PA VPNPYTPLSQILEEAEWFTVL
1			l			DLRDAFFCIPVHPDSQFLFAFED
1						PLNPMSQLTWTVLPQGFRDSPY
				i		LFGQALAQDLSQFSYLDTHVL
	l					WYMDDLLLAARSETLCHQATQ
						ALLNFLATCGYKVSKPKAQLC
1						LQQVKHLRLKLSKGTRALSEER
				i		IQPISAYPHPKTLKQLRMFLGIT
				i		GFCQIWINRYGKIARPLYTLIKE
1				1		TQKANTHLVRRTSLQPVAYLS
				1		KETGTSLQPVAYLSKEIDVVAK
						RWPHCLRVVAAIAVLVSKAVK
				l	1	MIQGRALTV WTSHDVNSILTAK
			1	I		GDLWLSDNHNEEKIEHNWQQV
			1	I		IAQTYAAQGDLLEVPLTDPDLN
				l		LYTDGSSFVEKGLRKAAYAVFS
1		ĺ	1	l		DNGILESNPLTPGTSTQLAELIA
		1			i	VTQALELEGKREPENPADNASY
1		1	1	1	l	SCEPLEDLRLLFRRQPIEAVKLQ
		1		ŀ		MVLQMEPQMQSMTKIYRGPLD
		1	1	1	l	WPASPCSNVDDIEGTPPKEISTA
$\overline{}$						

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14823	45191	A	14910	929	1324	RRYQVQRPSEGTLVYLFWPV*/ PVVGVDPWLPTGPD\YSLLVPIP PPRARTPRWA*A*MKGCRRSAT A*WCMC*SCRVPCPSCTPMFHS PCSAMTAVTPSLARP*THGSPP KAQGAPQGVKGPSSGLEAPPW A
14824	45192	Α	14911	2	286	
14825	45193	A	14912	220	249	P*N**PSTLTTGTGSTTT*QIRM KVN*MGN/DSPQGKNTPKMYS GEFSPVRVHVPFSLSDLKQIKID LGKFSDDPDGYIDVLQ
14826	45194	A	14913	1	163	EPENPADNASYSCEPLEDLRLL FRRQPIEAVKLQMVLQMEPQM QSMTKIYRGPLDWPASPCS*DF LGRGAFNVINIGAWASRPVQGS AVDLSHGLHLGLHLKNHL
14827	45195	Α	14914	1	287	
14828	45196	Α	14915	2498	2788	
14829	45197	Α	14916	112	333	
14830	45198	Α	14917	329	1666	
14831	45199	Α	14918	1	756	
14832	45200	A	14919	1256	1651	RRYQVQRPSEGTLYYLFWPV*/ PVVGVDPWLPTGPD\YSLLVPIP PPRARTPRWA*A*MKGCRRSAT A*WCMC*SCRVPCPSCTPMFHS PCSAMTAVTPSLARP*THGSPP KAQGAPQGVKGPSSGLEAPPW A
14833	45201	A	14920		987	MILVLGYNRKNTEGTYÖKÇKGT NASDFHIELSQVIEQVVSPKGSK EAQCCVLRHLGCESSESAPGIPP NLGIQLLTWANWUPFFTILA RAPSLALELMTQYFNNWNW YNNITDQGESKMSKLKGKERR KERKREKKEREKERELSQR KERKREKKGKKREREKERSQR LGQGYYYLTWRDIMLLLDQTL TPNERSAAITAVREFGOLWYILS QVDRKRIERERGPFTGQQAVP SVDPHWDTESEHGDWCHRYLL TDVLEGLEKTRKKRIN*SMISTI TOQKEENPINFLERLREALKH
14834	45202	Α	14921	473	1015	
14835	45203	Α	14922	3	937	
14836	45204	Α	14923	530	612	LQAKVFFLLRWPAILRG*NYGP RGVPH
14837	45205	A	14924	101	430	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
14838	45206	Α	14925	1	1587	MAALTLRGVRELLKRVDLATV
1				l		PRRHRYKKKWAATEPKFPAVR
						LALQNFDMTYSVQFGDLWPSIR
						VSLLSEQKYGALVNNFAAWDH
						PGDIVLDLCAAPGGKTLALLQT
1						GCCLTWSRKRCAALSMFSRPA
1				ŀ		RCLCRARTTAPRAVRRPDQRR
						ATAAATKQQARATPEAPGSAA
				-		HSSYCTMIPSACCCLPRQLKPPH
1				1		AQPARSAAKPGLDQPGSHARL
	İ					RGRGGEARQSLEAIRGGPSGIA
						PSDIPLPAKPSRRVIAQTYAARG
						NLLEVPLTDPDLNLYIDGSSFVE
İ						KRLRKAGIKAVKLQIVLQMEPQ
						MQPMTKIYCGPLDQPASPRSNV
1						DDIEGTNASDFPFLSQVLEQVV
						SPKGSKEAQCCVLRPLGCESNL
1						KQIKIDLGKFSDTPDGYIDVLQ
1	1					GLGQPYYLTWRDIMLLLDQTL
						TPNERSAAITAVREFGDLWYLS
	ĺ					QVNDRKITEEREQFPTGQQAVP
		1				SVDPHWDTESEHGDWCHRYLL
ì						TDVLEGLRKTRKKP\IN*SMISTI
L		_				TQGKEENPTAFLERLREALRKH
14839	45207	В	14926 14927	1	939	MOCHETIA WOODI DODI GROOV
14840	45208	Α	14927	l'	4317	MQSMTKMYCGPLDQPASPCSN
	1					VDDIEVKLEIMLLTERDDGKEV
		1				LDVGEERRYEIEEYDSKWTRK YNCLAVLRVHSGSOITTMKLIL
1		1				WYLVVALWCFFKDGKIAASRS
						GGFSYGSSSSGDLDRKKPLFSL
ı						EFGSPGETEDKSRQRQDAGSPK
						SEDTPAGGFFNSSSSSGDSDRTK
					*	PFFSLGLGAPGKAEDKSGDSOD
						AGGSKSEDTPPGGFFYGSSSSG
						DSDKKKPLFSFEFGATGEDEDK
						SRERWDAGNSRSEDS
14841	45209	В	14928	110	269	SIER W DAGNORSEDS
14842	45210	A	14929	1	561	EVFPACSQLVYWGKPITIYPLCE
17042	75210	ľ	14/2)	Ι'	50.	NNVYMLSPNASVCLYSP\LAEO
						FSHQFPSHDLPSVLAKFSLPVSL
	1	1		l		SEFRNPLAPAVQETQLIQMVV
	1					WMLQRRLLIQLHT\YVCLMASP
		l				SEEEPRPREDDVPFTARVGGRS
	1	l		l		LSTPNALSFGSPTSSDDMTLTSP
		l		1		SMDNSSAELLPSGDSPLNQRMT
	1	l		l		ENLLASLSEH
	L			L	l	DITECHOLOGIII

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
14843	45211	A	14930	1891	2522	GLSNQNQQPGRCIPDAHPVQDL
14643	43211	l ^A	14930	1091	2322	VODLVHTWCRVPCTSYVPCV*
		l				VPGAYLVHTRCIPGAGPGAYPV
		l				HTWCRTWCIPGAGFPAL/RPYH
		l				
l		l				ALLLLSDEKSLLGELPIDCSPAL
		l				VRVIKTTSAVKNLQQLAQDAD
						LALLQAMGSAVA\LVMWAWP
		l				RATGPLKYCLRVGAGGACGWR
						PRLVAVSRALLLVWRCGSPWG
		_				PALAWQKSPEYSTKVLEKQ
14844	45212	C	14931	113	274	
14845	45213	Α	14932	1	1634	MEIPAVILTVAPILHLTSCGFPV
						TGDGLKQAEGKCAPEEWNPTK
		1				FIKLTDNQACFACGLWPLSEIQ
						RRTSRTVTIISRISALREKGNGR
		l				KPTSISGSVNEMRQVLVKDQVS
		l				MPRVLPTIIFVPDEAGFSAGAV
						WPCAAPLWGSSWTPLKVENCS
						FYIRLSFAISVGGDELKIAKLVS
						AFPERAYDLVGHQHQRPKVDK
						TTKMGRNQSRKAENSKNQNMS
						SPPKEHNSSPAREQNWMENEF
				1		DELREVGFRRSVITNFSELKEH
						VLTHHKEAKNLEKRLDEWLTR
		l				INSVEKSLNDLMELKTTLRELR
						EAYISFNSQFDQAEERVSVIEDQ
						ISEIKQEVKFREKRVKRNEQSL
						QEIWDYVKRP\NLRLIGIPESDE
				1		ENGTKLENTLQDIIQENFPNIAR
						QANIQI/QGKLQRTPTKVASSRT
						ATPRHIISVGSPKLNMK\EKILR
		l	1			AAQRERSGLPTKGKPFRLTAAD
1		l	1			LSAETSTSQKTEWGPIFNILKEK
		l				NF\QPR\ISYPAK\LSFISEGEIKYF
			1	l		TDQAKCLRDF\VTTRAFSCLP*K
1		1	1			STRPALKELLKGST
				L		3 I KT ALKELLKUS I

NO:	No. of peptide sequence sequence codon for peptide codon	SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
14846 45214 A 14933 T 1429 MVLAAGDPVEKGITWARENA TVISCTMMKLSMIPIGDQSYR IPLKFVACILLEGNARCWPLPE DPSOJRILIVYQDCERGRNVL DSSVRRNEDISVSRIKYLGJQ TRDVKDLFKENYKPLINEIKE TNKWKNIPCSWERNIVKAIII KKLGSDAQVKVFGKCCQLKP GDSSSLDSSVTSSSSKIRQQCT TRDVKDLFKENYKPLINEIKE TNKWKNIPCSWERNIVKAIII KKLGSDAQVKVFGKCCQLKP GDSSSLDSSVTSSSSKIRQQCT TRDVKDLFKENYKPLINEIKE TNKWKNIPCSWERNIVKAIII KKLGSDAQVKVFGKCCQLKP GDSSSLDSSSLTSSSTSDIRDQCL YQRTVDLSAQLSSSGAGQTT SGSLTPMTPHWEIPPSGQQT HTGELRLASGRGSSGMKLPEE TDRNLCCSAASSGDTQANNED SGPPANTSRPAAGPDCDREN TKLENTLADIHENFPNIARQA IKIQEIQRTQQRYCSRATPRH VRFTKVEMKEKMLRAAGEKK WVIHKGKSIRLTVDLSABTLQ RREWGPIPNVLKEKNPQLRIS PAKLSFISEGEMKSFPEKQMLI DFVTIRLAQELLKEALNMAR NRYQPLQKHM TRISTALQELLKEALNMAR NRYQPLQKHMAR TRISTALQELLKEALNMAR NRYQPLQKHKNLEASHYLTI. TRISTALQELLKEALNMAR TRISTALQELLK	14846 45214 A 14933 I 1429 MVLAAGDPVEKGITWARENA I 1429 MVLAAGDPVEKGITWARENA I 1429 MVLAAGDPVEKGITWARENA I 1429 MVLAAGDPVEKGITWARENA I I 1429 MVLAAGDPVEKGITWARENA I I I 1429 MVLAAGDPVEKGITWARENA I I I I I I I I I							
14846 45214 A 14933 I 1429 MVLAAGDPVEKGITWARENA IVISCTMMKLSMIPIGDQSYR IPLKFVACILIEGNARCWPLPE DPSOIRLIVYQDCERGRONUL DSSVKRRNEDISVSRIKYLGIQ TRDVKDLFKENYKPLINEIKE TNKWKNIPCSWERNIVKAIII KKLGSDAQVKVFGKCCQLKP GDSSSLDSSVTSSSDIKDQCL QQRTVDLSAQLSSGGGGTTS SGSLTPMTPHWEIPPSGGQCT HTGELRLASGRSGMKLPEE TDRNLCCSAASSGDTQANRD SGPPANTSRPAAEGPCDCREN TKLENTLKDIIHENFPNIARQA IKQEQRTQGRYCSRRATPRH VRFTKVEMKEKRMEN, AEAGEPCD REWGPIPNVLKEKNPQLKB PAKLSFISEGEMKSFPEKQMLI DFVTIRLAQELLKEALNMAR NRYQPLQKHM NRYQPLAKHADA NRYQPLQKHM NRYQPLAKHADA NRYQP	14846 45214 A 14933 I 1429 MVLAAGDPVEKGITWARENA INISCTMMKLSMIPIGDQSYRN IPILKFVACILIEGNAKCWPLPE DPSQIRLIVYQDCERGRANUL DSSVKRRNEDISVSRIKYLGIQ TRDVKDLFKENY FELLINEE TINK WKNIPCSWIERINIVKIAII KKLGSDAQVK VFGKCCQLKP GDSSSSLDSSVTSSSDIKDQCL YQRTVDLSAQLSSSGKGQTTS SGSLTMTPTWEIPPSGQQTS HTGELRLASGRGSSGMKLPEE TDRNLCCSAASSGDTQANRDV SGPANTSRPAAEGPDCDREN TKLENTLKDIHENFPAIAQA IKIQEIQRTQQYCSRRATPRE VRFTKVEMKEEKMLRAAGEK- WVIHKGKSIRLTVDLSAELQ RREWGPIPNVLKEKNPQLRIS PAKLSFISEGEMKSFPEKQMLI DFVTIRLAQELKEALNMAN NRYQPQLQKHM NRYQPQLQKHM FSKLTRGVKDLFRENYKFPLLN IKRGKQMEEHSMLMGRKNQ YREKWYDLFVNL-1QCPHPIQ TNDCPHRIGKALL*SSYGTKK PALPSQS*DKRTLEASHYLTS NYTTRLQ*PRQHYWONDLD DQWNRTEASEIMPHIYTILLW NLRKTSMGERIP 14848 45216 A 14935 2 872 14936 14937 14936 14937 14938 1 14938 1 14937 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14938 1 14940 235 337 QQAHKTPQRRHSAFTEYVRN RLPETVIRVQPALAIFISPDIAF FVHFRADDDVTARVHIAIMIE WCDR VEKRVSDA VACFTLESI AEVGVLCRTHIRQ*RQVTPAR *ALSRILHACRSEDVENNREPP VIRVQPALRIFIAGPTARVHAIMIE WCDR VEKRVSDA VACFTLESI AEVGVLCRTHIRQ*RQVTPAR *ALSRILHACRSEDVENNREPP VIRVQPALRIFIAGPTARVHAIMIE WCDR VEKRVSDA VACFTLESI AEVGVLCRTHIRQ*RQVTPAR *ALSRILHACRSEDVENNREPP VIRVQPALRIFIAGPTARVTPARVENDAR VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVRN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTEYVEN *ALSRILHACRSEDVENNREPP VIRVQPARRISAFTE							deletion, \-possible nucleotide insertion)
IMSCTMMKLSMIPIGIGGSYR IPLKFVACILIEGNAKCWPLPE DPSQIRLIVYQDCERGRNVL DSSVKRRNEDISVSRIKYLGIQ TRDVKDLFKENYKPLLNEIRE TNK WKNIPCSWIERINIVKIAII KKLGSDAQWKPGKCOLKP GDSSSSLDSSVTSSSDIKDQCL YQRTVDLSAQLSSSGKGQTTS SGSLTPMTPHWEIPPSGQQTT HTGELRLASGRGSSGMKLPEE TDRNLCSAASSGTOANED SGPPANTSRPAAEGPDCDREN TKLENTLKDIHIENFPNIARQA IKIQEIQRTQQRYCSRATPHEI VRTFKVEMKERMLRAAGEK WVIHKGKSIRLTVDLSAETLQ RREWGPIENVLKEKNPQLRIS PAKLSFISEGEMKSFPEKQML DFVTIRLAQELLKEALNMAR NRVQPLQKHM NRVQPLQKHM NRVQPLQKHM SSLTIRGVKDLFRENYKPLLN IKRGHKQMEEHSMLMGKNN YREKWYLPKVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNL IQCIPHEI TNDCFHRIGKKLL*SSYGTKK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVNC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVC INCHEI TNDCFHRIGKKLL*SSYGTK PALPSQS*DKRTKLEASHYLT: NYTTRLQ*PRGHYNVC INCHEI TNDCFHRIGKKLL*SSYGTK TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE TNCFHRICTOLSTAGE	INSCTMMKLSMIPIGGGSYRN PILKFVACILIEGNAKCWPLPE					sequence		
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*ALSRILHACRSEDVRHNRPPY VIRVKQPAIRRDLAPT*PHCSS AQTITSLPGCMREVI/ELRPEF K*RKTVLRPTPIMRARAERRC	*ALSKILHACRSEDVRHNRPPY VIRVKQPARRDLAPT*PHCSS AQTITSLPGCMREVT/ELRPEP! K*RKTVLRPTPIMRARAERRCI 14854 45222 C 14941 I 1755 14855 45223 A 14942 I 747		l	1		1	1	
VIRVKQPA\RRDLAPT*PHCSS AQTITSLPGCMREVI/ELRPEF K*RKTVLRPTPIMRARAERRC	VIRVKQPA\RRDLAPT*PHCSS AQTITSLPGCMREVI7ELRPEFI K*RKTVLRPTPIMRARAERRCI 14854 45222 C 14941 1 1755 14855 45223 A 14942 I 747	1	1	1			1	
AQTITSLPGCMREV1/ELRPEF K*RKTVLRPTPIMRARAERRC	AQTITSLPGCMREVI/ELRPEF K*RKTVLRPTPIMRARAERRC L4854 45222 C 14941 1 1755 L4855 45223 A 14942 1 747	1	1	1			1	
K*RKTVLRPTPIMRARAERRC 14854 45222 C 14941 1 1755	K*RKTVLRPTPIMRARAERRCI 14854 45222	l	1	1		1		
14854 45222 C 14941 I 1755	14854 45222 C 14941 1 1755 14855 45223 A 14942 I 747	1	ŀ	1			1	
	14855 45223 A 14942 I 747							K*RKTVLRPTPIMRARAERRCK
14855 45223 A 14942 1	11000 11000			-				
11000 11000	14856 45224 B 14943 I 12466						1	
14856 45224 B 14943 I 2466	1-	14856	45224	В	14943	1	2466	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14857	45225	А	14944		333	LNLVLFQQRAELMIS*PQ*FSRA LLVIFPLLQTELQQLFFLLHVNP AAVEAPPSVPIGLYSENVRNNR LPETVIRVKQPALARFSPDIAPL FVHFRADDDVTARRVEKRCK
14858	45226	В	14945	1	1794	
14859	45227	A	14946	1183	1899	LINRRDSYVEYLADKVADREED AEFYLEAIMEARYTVAGMGLV MEVQDYFDGEADRLAKAWLA EYTPQIKSLKDERKEA YRQIVE MSTEPQDVDLVRPANKFEMTR VREGEKEADLPVWKHHLJSVY RKELSGSVEPLONOVFEIETKRE NS*EL**KSLKDERKEAYRQIVE MSTEPQDVDLVRPANKFEMTR VREGEKEADLPVWKHLLCDE SGNYPALLNHWETRFLRSKPN VKHRSSOKSPYTFVFFISQYDL PAFYPINGHFLVERVVKVRIVFI EEFADHHILAIPCNNRLQRHA
14860	45228	В	14947	1	3099	
14861	45229	A	14948	1174	1545	
14862	45230	Α	14949	1	945	
14863 14864	45231 45232	A	14950	62 1	576 1642	MLANIGFYNSNVIHRSKGAIGE GPPDRFGIVEQNLGGIDSIPCYN GPPRRSAQKTTLGFLNOPRIPS AQCAGAFPRVAAAGAGGGG GGSGGGCLGWELRRRQRR RRWRRGEGEGDODAGHMEPR PAGRAEARRRRLRGNPPGAR EPNERGNGSGYSWAAEVSODG RGARTEAARTRRPGGADNYP GA VPPVQLLTGWCVTAKAPPDI SAISALTAVKHGNCSSLTPLLNP PGSDVIVCAEMDEQWGYVGAK SRQRWLFTAYDSLRKTVVAHV FGERTMATLGRLMSLLSPFDVV IMMTDGWPLYESRLKGKLHVI IMMTDGWPLYESRLKGKLHVI TRENVESDYSWVENERFGELR HPVDEEKARCLEGIGGHTRGLV ASLDMCLEGGGGTGGLAQAE LCWNSSEMRTTMSSSGSSTPWP PVSRRRPWTERRLGSVRGDRR ERSRRAVNRTPEMSRVYSGAPR GTESRRWWMMKCWDHQMRG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
14865	45233	Α	14952	1	1513	MIIALQKESLMSLQAQLDKALQ
						KEKHYLQTTITKEAYDALSRKS
						AACQDDLTQALEKRTGRDWNP
						PPFRTQQRGSKSLTAIRPLCLFR
						KLNHVTSETKSLQQSLTQTQEK
						KAQLEEEIIAYEERMKKLNTEL
						RKLRGFHQESELEVILYEEEMG
				i		NHNENTGEKLHLAQEQLALAG
						DKITSLERSLNLYRDKYQSSLS
						NIELLECQVKMLQGELGGIMG
ļ						QLSKTSSDLSKM/RCY/TTW*M
						WTKLQFIHSILTSVIFHLLSFSSM
				1	i	GSI*KWIMALSGSSFLLDAHYIS
						QFGPRQLPSVCAEPEKEKATDG
						VVRNGKSTAGHQRYLCSHCRK
						TWQLQFTYTASQPAGQ*RHRL
						RG\MDEQWGYVGAKSRORWLF
						YAYDSLRKTVVAHHREQGSIK
			ļ			CKLEEDLQEATKLLEDKREQLK
	İ					KSKEHEKLMEGELEALRQEFK
l						KKDKTLKENSRKLEEENENLR
						AELQCCSTQLESSLNKYNTSQQ
						VIQDLNKSPMEIRFQLYIFTPDQ
14866	45234	Α	14953	125	363	INVSVWSQPFPGG*QPEKGLKE
		l				HPADARPGSC\PPAGRPVQGRA
		1				EVASAAPPRGSSPAAAPGQSHP
						PSCLRAPLYWSFSR
14867	45235	A	14954	306	583	TCSTRRRKNDLSVKYVSVWSQ
		l				PFPGG*QPEKGLKEHPADARPG
		ŀ				SC\PPAGRPVQGRAEVASAAPP
		ŀ				RGSSPAAAPGQSHPPSCLRAPL
						YWSFSS
14868	45236	Α	14955	225	407	LRGMVTFSPFSSILNGQFEPNSG
	1					CPL/CPVCWVCLREPGDPEKLG
						EFLQKDNISVHYFCLVS
14869	45237	Α	14956	484	722	NTCRCGHNRSPEVNSRRGGLRS
l		1	1	l		IPRMPDAGSCS/PPAGRPSAGPG
l		1				RRSPPPLLPRGSSPAAAPGQSHP
l	1	1			1	PSSSOGAAVLEL

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i		sequence		
	14000	<u> </u>	14055	1	1,700	CAMERDACTIO A CHOICE ON THE CAMERON IN
14870	45238	A	14957	434	1790	GNFFRKTISACIISVLSYLVSCLR
						GASPTEASMDFCLKTSKREAAR
1						ASRKICFVCKKKGAAINCQKDQ
						CLRNFHLPCGQERGLPFTIFWR
1						VQIIL*QTSP/EHRTSNMGHVGE
						ESCILCCEDLSPTECLRTFQSPC
					1	CSQAIYHRKCIQ/TNMPTHQSK
						HFFKCPQCNNRKEFPQEM/REN
						GNSYSRQRCCLGTRARGFLRLI
				İ		SALSAL*CPHLVCMNKAETALR
						MKGGGASFCVLHADPTEPTGT
						APLLDLTVRNGSVRSVHL/PAA
		l				TDYIPENSGDIPCCSSTFHPEEHF
						CRDNTLEENPGLSWTDWPEPSL
1						LEKPESSRGRRS/VLLEVQGCQ
						NH*QLQKIQVTPSE*LLVPHNR
						V*TCAPPSGLGREHSGTVRQGS
		İ				RASSETMS/LSKEKSQWSMRRE
		1				OSRCOOGNAFMATRVPLLSPPL
l		1	İ			LPPRILPP*SCFHMPLLTSPMFV
İ	ĺ	1				VMQIKVFSPKKKKK
14871	45239	A	14958	146	484	
14872	45240	A	14959	228	353	
14873	45241	A	14960	214	722	TARELPGRVLKEMYIOOK*EGA
14075	13211	ľ.,	1,1,00	1	/	ALGRPTRPGRQGQRVRRAECR
						RLFRGVLRPCNRC
14874	45242	В	14961	1	951	Ida no value cinte
14875	45243	Ā	14962	241	517	QSCGQRLPTVLR/PLRTPWILPM
14075	7,52,73	(`	14702	1-11	J***	YPOPLPAGGPTPCLRSRPYINITI
		1				LKGDKGDPGPMGLPGYMGRE
		l				GPOGEPGP*GSKGDKGEMGRP
		1				RRPVP
14076	45244	-	14062	748	1000	
14876	45244	Α	14963	/48	1055	DRKQTSPGDAYHTIPGSLSSSSD
		1	1			HVLSEPD*GEGSHL**SVQHDD
		1				VPPTA**RLQY\RAVVLFPTATL
		l		1		PAILIM*GMRCPLVPRNCSRTW
				l		TAQIRAYIEVQQT

K=Unknown,
e nucleotide
eotide inscrtion)
AAVLPPDGA
FCHHGGVHI
IDRVGQIPIPE
PPSRHTEDRE
NRWFLPVPT
RTCSMQGLSF
DLRGNVSKES
FPMEGGQKD
NLSGPHGPGA
SEISLSVGQV
SSHCRGARL
QDEFSLQPEE
LEQYVVVA/
VGQ/VVDIIE
AEEQGWVPA
FLCSLKKML
IQIKTTMRYH
NNRCWRGC
CKLVQPLWK
PFDPAIPLL
NSHITILTLN
ASWIKSQDP
DTHRLKIKG
KAGVAILVS
KEGHYMMV
MYAPNTGAP
LDSHTLIMG
RQKVNKDTQ
IYRTLHPKST
KIDHILGSEA
'LSDHSAIKL
ŀ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14887	45255	Α	14974	1	1240	MNIDAKILNKILANRIQQHIKKL
		1				IHHDQVGFIPGMQGWFNICKSI
		l				NVIQHINRAKDKNHMIISIDAEK
		l				AFDKIQQPFMLKTLNKLGIDGM
		1				YFKIIRAIYDKPTANIILNGQKLE
		l				AFPLKTGTRQGCPLSPLLFSIVL
		l				EVLARAIRQEKEIKGIQLGKKE
		1				VKLSLFADDMIVYLENPIVSAQ
		1				NLLKLISNFSKVSGYKINVQKS
		1				QAFLYTNNRQTESQIMSELPFTI
		l				ASKRLKYLGIQLTRDVKDLFKE
	ĺ	1				NYKPLLKEIKEDTNKWKNIPCS
		l				WVGRINIMKMAILPKVIYRFNA
		l				IPIKLPMPFFTELEKTTLKFIWNE
		1				KRARIAKSIPSQKNKAGGITLPD
		ı				FKLYYKATVTKTAWYWYQNR
		ı				DIDQWNRTEPSEITPHIYNYLIF
		l				DKPEKNKQIQKAFILH\KEPWV
		_				MWNMGTNVR
14888	45256	A	14975	1	2241	
14889	45257	Α	14976	11	2802	MYGTGYRDVAGKWAIDPDSK
		l				REFLDVTQEGIQGCDFSGTCRQ
		l	1			TLSILTQPLRQWGLEGIKKPNS
		l		1		WIISEESVSNGGPPLLIPRQTAS
		l	ļ	ł		GVDLQQTPTDLQLRVLTVRRK
		l	1	ĺ		TNKQKGIASTSTKRTSTPKPHL
		l				YVTIIKDQSYIKPQRWGKNIAE
		l				KLKILKIRVALSLQRNAAPHQQ
		1		l		WNKAGRRMSLMSSQKKASEVI
				l		ESQMNEIKGEEKFREKRVKRNE
						QSLQEIWDYVKRPDLRLIGVPD SDGENGTRLENTLODI
14890	45258	A	14977	1	2739	SEGENGT REENTEQUI
14891	45259	A	14978	i	2478	
14892	45260	A	14979	590	727	
14893	45261	A	14980	1	3255	
14894	45262	A	14981	1	3810	
14895	45263	В	14982	i	3127	
14896	45264	A	14983	i	3325	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14898	45266	A	14985		2067	MELKTKARELREECRSLRSRCD QLEERVSAMEDENNEMKREG QLEERVSAMEDENNEMKREG KFREKRIKRNEGSLQEIBUDYYK RPNLHLIGVPETDGENGTKLEN TLODHIGDENFDLARQANVOIGE IQRTPQRYSSRRATPRIHIVRFT KVEMKEKMLRAAREKEIQTTIR EYYKHLHANKLENLEFMDKFL DTYTLPRLNGEKVESLNRPITG AEIVAINNSLEPKESPOFDGFTA EFYQRGILPNSFYEASIILIPKLG ROTAKKENFRISLMNIDAKILN KLAKRIQQHIKKLIHHDQVGFI PGMQGWFNIRKSINVIQHVNRT KONNHMISIDEEKAFDKIQDP MLKTLNKLGIDGMYFIIRAIVD KRYAHILNGKKLEAFDLKTGM MIYULENPIVSAQNLLKLISNFS KVSGYKINVOKSOTFLYTNNR QTESQIMSELPFITASKRIKYLGI QTENDVKDLFKENYKRVLKEIK EDTNKWKNIPCSWVGRNIVK MALIKKVIYRFNAIPIKVPMTFF TELEELTTLKFIWNGKRACIAKSI LSQKNKAGGITLPDFKLYYKAT TELECTTILKFIWNGKRACIAKSI LSQKNKAGGITLPDFKLYYKAT FESEITTHIYNYLIFDKRACIAKSI LSQKNKAGGITLPDFKLYYKAT EFSEITTHIYNYLIFDKRACIAKSI LSQKNKAGGITLPDFKLYYKAT FESEITTHIYNYLIFDKREKNKYLGI VGKOSLFFHYNYLIFDKREVKYKAT FESEITTHIYNYLIFDKREKNKYLGI VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREKNKY VGKOSLFFHYNYLIFDKREVKNY VGKOSLFFHYNYLIFDKREVKNY VGKOSLFFHYNYKWCWENNULAICR
		_				KLKLDPFLTPYTKINSRWIKD
14899	45267	A	14986	1	1293	
14900	45268	Α	14987	1	3229	
14901	45269	Α	14988	1	3352	
14902	45270	Α	14989	1	1419	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
14903	45271	A	14990	1	1716	MKAEIKMFFETNENKDTTYQN LWDTFKAYCRGKFIALNAHNR KQFRSKIDTLTSQLKELEKQEQ TYSKASRRQEITKIRAELKEIET QKTLQKINESSWFFERNIKIDR PLARLIKKKRQKNQMDTIKND KGDITTDPTEIQTTYREYYKHL LNQEEVESLNRPITGAEIVAIINS LPTKKSPGPDGFTAEFYQRYKE ELPGRDTTKEENFRISLIMNIDA KILNKILANQIQQHIKKIIHHDQ VGFIPGMGQWFNIRKSINVIQH NRAKDKNHMIMSIDAEKAFDKI QQFFMLKTLNKLVLEVLARAIR REKEIKGIQGEVEKLSLFAD HMIVYLENPIVSAQNLKLILSING QTESQIMSELPFTIASKRIKYLGI QLTRDVKDLFRENYKPLLKEIK EDTINKWKNIPCSWYGRINIVKV AILPKYLYSFNAIPKIPCSWGRINIVKV AILPKYLYSFNAIPKIPCSWGRINIVKV AILPKYLYSFNAIPKIPMFFFTEL EKTILKFIWNQKRAPKLYFIVTK KNRAGAITLPDFKLYYKPTVTK TAWYSYQNRDIDQWNRTEPSE
14904	45272	Α	14991	1	702	

NO: of p		hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14905 45	2273	Α	14992	473	2385	HISPLISTIVLMYLTMDLTRSQKI GWASSPSHGTSTPWPYRLR TAG KPCKWQVSLCWWLSASPGSSF SLVRSSSLLIIVPHAESVSRRPSLL KPCKWQVSLCWWLSASPGSSF SLVRSSSLLIIVPHAESVSRRPSLL KVGLFPARY*HEVWQRSPSPQ TRDHLIKRNPSQDV*RQQPRG TRJHLIKRNPSQDV*RQQPRG TRSYNBKCSLHTSALGIGRQPSC DATL*GQQGLPVK*KRQQPAP MSQQQTGAALEVLARAIRQEK EKGGIQ.GREEVKLSLFADDMI VYLENPTVSAQNLLKLISNFSK SQYKINVQKSQAFLYTNIRQ TESQIMSELPFTIASKRITYLGIQ LTRDAKDLFKENYKPLLKEIKE DTNKWKNIPCSWVGRINIVKM ALLPKEPGWARSLVFGARGLQG QGKCGQSRGGGSGCLGGGGGST ENWQHKEPSARKTGSTENRH REPAAPRTGSTENQHREPAR TGTENRHEPSARKTGSTENRH REPAAPRTGSTENQHREPAR TGTENRHEPSARKTGSTENRH REPAAPRTGTENRHEPSAP TGTSTENRQHREPARTGTENRH REPAAPRTGTENRHEPSAP RTGSTENRQHREPARTSTEN RQHRESAAPRTGTENRHEPSAP RTGSTENRQHREPARTSTEN RQHRESAAPRTGTENRHEPSAP RTGSTENRQHREPARTSTENRH RPARHELVETTENRHEPSAP RTGSTENRQHREPARTSTENRH RPARHELVETTENRHEPSAP RTGSTENRGHREPARTSTENRH RPARHELVETTENRHEPSAP RTGTSTENRGHREPARTSTENRH RPARHETUTENRTGTENRRPEP PAGTREPRHTPMRGGSEGALFP PAGTREPHTPMRGGSEGALFP PAGTREPHTPMRGGGEGGGF PPYA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=pnssible nucleotide deletion, \=possible nucleotide insertion)
14906	45274	A	14993	687	2274	LRDCKRINQHPVKTDQSAFCK M/API/MQDVVLEVLARAIRQEE EIKGIQLGKEEVKLSLFADDMI VYLENPIVSAQNLLKLISNFSK SQYKINAQKSQAFLYTNNRQTE SQIMSELPFTIASKRIKYLGIQLT RDVKDLFKENYKPLLKEITEDT KWENIPGSWYGRNIVKMAIL PKVIYRFNAPIPKLPMIFFTELE KTTLKFIWNQKRARMAKSILSC KNRAGGITLPDFKLYYKATVI KTAWYWYQNRDVDQWNRTEE SEITPHIYNYLIFDKPDKNKQW GKDSLENKWGWENWLAICRKI KLDPFLTPYRINSRWKIDLNV RPKTIKTLEENLDHTIQDIGMGK DFMSKTPKAMATKAKIDKWDI EIKLSFCIAKETTIRVNRQPTKW EKIFATYSPDKGLISRIYYELKO KJEKSFCIAKETTIRVNRQPTKW EKIFATYSPDKGLISRIYYELKO KKKTINNFKWAKDMINRIFF KEDIYAAKKHMKRCSSSLAIRE IQIKTTMRYJILTPVRMAIKKSC BNJRCFCWGPOSPG-ALRSWF EDIAGTSYVGMNQCQMRKNTT TLI
14907	45275	Α	14994	1	3189	1.22
14908	45276	A	14995	1	3351	MGKKQNRKTGNSKMQSASPPK KERSSPATDQSWMENDFDEL REGGFRRSNYSELREDIGTKGK EVENFEKNLEECITRITNTEKCL KELMELKTKARELREECRSLRS RCDQLEERVSAMEDEMNEMK QEGKIKRDKEGHYIMYKGSIQQ EELTILINIYAPHTGAPRFIKQVI. SDLQRDLDSHTLIMGDFNTSLL TLDRSMRQKVNKDTOELNSSL HQADLDIYBTLIPKSIEYTFS APHIHTYSKIDHIVGS
14909	45277	Α	14996	1	3192	
14910	45278	Α	14997	1	2742	
14911	45279	В	14998	1	2313	
14912	45280	Α	14999	1	1542	
14913	45281	Α	15000	1	3300	
14914	45282	Α	15001	1	1824	
14915	45283	Α	15002	I	2982	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14916	45284	Α	15003	li .	2375	MSGFLISPKHKHGTNSGIVSWA
		1				DVQSQLRHIEAAFDQEAAAVL
	1	1				MARLGVFRAESEEGPDVLRWL
	1	l		1		DRQLIRLHSLLTTLDIVPAGKGE
i		l				VVKGPMSISQQAKGCQKFGQY
						NKEDPTSFRLSDSFSLYPQFMF
						HLRRSPFLQVFNNSPDESSYYR
						HHFARQDLTQSLIMIQPILYSYS
						FHGPPEPVLLDSSSILADRILLM
		1				DTFFQIVIYLGETIAQWRKAGY
						QDMPEYENFKHLLQAPLDDAQ
	ì	1				EILQARFPMPRYINTEHGGSQK
İ		l				KREKNQIDAIKNDKGDITTEPTI
		l				IQTTIREYYKHLYANKLESLEE
l		1				MNKFLDTYTLPRLNQEEVESLN
	1	1		l		RQITGSEIEAIINSLPTKKSPGPD
	1	1				RFTAKFYQRTNDQKHMIISIDA
l	1	l				EKAFDKIQQPFMLKTLNKLGID
		l				GTYFKIIRAIYDKPTANIILNGQ
	1	l				KLEAFPLKTGTRQGCPLSLLLF
	l .	l				NIMLEVLARAIRQEKEIKGIQFG
		l				KEEVKLSLFADDMIVYLENPIV
	i					LAQNLLKLISNFSKVSGYKINV
	1	1				QKSQAFLYTSNRQTESQIMSEL
						PFTIASKRIKYLGIQLTRDVKDL
					l	FKENYKPLLNEIKEDTNKWKNI
l	1	1			1	PCSWVGRINIVKMAILPKVIYRF
			1		ł	NAIPIKLSMTFFTELEKTTLEFI
	1	1			[WNQKRARITKSILSQKNKAEGI
	1	1			1	TLPDFKLQYKATVTKTAWYSY
	1	1		ł	1	QNRDIDQWNRTEPSEIVPCIYN
						YLIFDKPDKNKKWGKDSLFNK
14917	45285	Α	15004	I	1428	
14918	45286	Α	15005	2	1975	

NO: of peptide sequence bod in USSN 109/549,217 location of first codon for peptide sequence of peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion, peptide deletion peptide deletion. Peptide deletion peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide deletion. Peptide de	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
14919 45287 A 15006 I 3891 MKPEIKMFFETNENKDTTYQ WDTFKAVCRGKFIALNAHKR QERSKINITING/FKELEKQEQ NSKASRRQEITKIRAELEKEIET KPLQKINESSRYFEKINKIDE LARLIKKREKENGIDAIKNDK DITTDPTEIQTTIREYYKHLVA KLENLEEMDKFLDMYTLPEL QEEVESLMRPITGSELBINSL TKKSRQPDGFTAEFYQRYKEE VSFLLKLFQSIKKEGILLNLFY DSIILKRPGRGDTYKKENRFPE MNIDAKILKILANRIQQHIKK HHDQVGFIFGMQGWFNIRKSI VIQHINRTKOKNHMISIDAEK FDKIQQFFMLKTLNKLGHGT LKHRAYDKPANILINGQKL AFPLKTGTRQCPLSPLVFNI EVLAWAHQKEKEIGGLGKE VKLSJFADDMYVLENPIISAN NLLKLISNFSKVSGYKINVOK: QAFLYTNNOGTSLIMSELPF ASKRIKYLGQLTROVKOLFK NYKPLLNEIGDTSLIMSELPF ASKRIKYLGQLTROVKOLFK NYKPLLNEIGDTSLIMSELPF ASKRIKYLGQLTROVKOLFK NYKPLLNEIGDTSLIMSELPF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKAGGTITLF FKLYYTATVTKTAWYWYON MVLVPKQRYKSMEQNKALGEL DEQQRKRLEELFLTYQKGVCG KDDDFEKISELGAGNGGVFF VSHKPSGLVMARKLHEIEP, RNOINELGVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCFGFLWMENSDYGLLLLPKNI KARNSHIGDVVVVELLPKNI WKGRTVALCENDC							
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DSIILERKPGROTTIKERNERPEN MNIDAKILIKILANRIQQHIKK HHIDQVGFIPGMQGWFNIRKSI VIQHINRITKOKNHMIISIDAEK FIDKIQOPFMLKTIKNLGHGHG LKIIRAIYDKPTANIILNGQKLI AFPLKTGTRQGCPLSPLVFNIV EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISA NLLKLISNFSKVSGYKINVQK QAFLYTNNGQTESLIMSELPF ASSRIKYLGIQLTRDVKDLFK NYKPLLNEIKEDTTNKWNIP WVGRINIVKMAILPKUYBEN PIKLPMTFFTLELEKTTLNFIWN KRARITKSILSQKNKAGGITLF FKLYYTATVTKTAWYWYON MVLYPKQRYRSMEONALRIK NAACLQLSDL MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRRELBEATTOKOKVGE KDDDFEKISELGAGNGGVFF VSHKPGLVMARKLHELEE, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLDL LCLSFISSKVISDLVSDLIHGI KARNRSHIGDVVVVELLPKNI WKGRTVALCENDC			1				TKKSPGPDGFTAEFYQRYKEEL
MNIDAKILIKILANRIQOJHIKK HHDQVGFIPGMQGWFNIRKSI VIQHINRTKDKNHMIISIDAEK FDKIQQPFMLKTLNKLGHGT LKIIRAIYDKPTANILINGQKLI AFPLKTGTRQCGPLSPLVFNI EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISA NLLKLISNFSKVSGYKINVOK: QAFLYTNNOGTSLIMSELPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHEICDTTSLWSLEPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHEICDTTSLWSLEPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHEICDTTSLWSLEPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHEICDTTSLWSLEPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHEICDTTSLWSLEPF ASKRIKYLGIQLTRDVKDLFK NYKPLLISHENDTTSLWSLEPT WVGRNIVKMAILPKVIYRFN PIKLPMTFFTDLEKTTLINFIWN KRARITKSILSQKNKAGGITLEF FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNKALGITLEF FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNKALGITLE FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNKALGITLE FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNKALGITLE FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNKALGITLE FKLYYTATVTKTAWYWYON WYGRATINLEALQKKLEELE DEQQRKRLEAFLTQKQKVGG KDDDFEKISELGAGNGGVVFF VSHKPGLUMARKLIHLEIKP, RNOJIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCFGFLWMENSDYGLLLLDLL LCLSFSTSKWISDLVSDILHGI KARNSHINGDVVVVSDLJKNI WKGRTVALGENDC	1				ľ		VSFLLKLFQSIKKEGILLNLFYE
HHDQVGFIPGMQGWFNIRKSI VIQHINRIKDKNHMIISIDAEK FDKIQQPFMLKTLNKLGHLGT LKIIRAIYDKPTANILINGGKLI AFPLKTGTRQGCPLSPLVFNI) EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISAI NILLKLISNFSKVSGYKINVQKI QAFLYTNNGOTESLIMSELPF ASKRIKYLGIQLTROVEDLFK NYKPLLNEIKEDITNKWKNIP WVGRINIVKMAILPKVIYRFN PIKLPMTFFTLDLEKTLINFIW KRARITKSILSQKNKAGGITLE FKLYYTATVTKTAWYWYQN MVLVPKQRYRSMEQNRALRS NAACLQLSDL H4920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWIRTNLEALQKKLEELE DEQQRKRLEAFLTQKQKVGE KDDDFFKISFLGAGNGGVFF VSHKPSGLVMARKLHELER, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFNFSKVISDLVSDLHGI KARNSHIGDVVVVELLPKNI WKGRTVALGENDC					l		DSIILKPKPGRDTTKKENFRPISL
I VIOHINRTKOKNHMIISIDAEK FDKIQOPPMLKTINKLGIHGT LKIIRAIYDKPTANIILINGQKLI AFPLKTGTRQCCPLSPLVFNI EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISA NLLKLISNFSKVSGYKINVOK. QAFLYTNNGQTESLIMSELPF ASKRIKYLGIQLTRDVKDLFK NYKPLLNEIKEDVITNKWKNIP WVGRINVKMAILPKYIYRFN PIKLPMTFFTDLEKTILNFIVN KRARITKSILSQKNKAGGITLE FKLYYTATVTKTAWYWYON MVLYPKQRYRMEQNALARI NAACLQLSDL 14920 45288 A 15007 1 4215 MHSCHDATGGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEEATTOKQKVGE KDDDFEKISELGAGNGGVFF VSHKPSGLVMARKLIHLEIKP, RNOJIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFSFSKVISDLVSDLIHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC			ļ.		İ		MNIDAKILIKILANRIQQHIKKLI
FDKIQQPFMLKTLNKLGHGT LKIIRAIYDKPTANIILNQGKLI AFPLKTGTRQGCPLSPLVFNIII EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISA NILLKLISNFSKVSGYKINVQK. QAFLYTNNQOTESLIMSELPF ASKRIKYLGQUTRDVKDLFK NYKPLLNEIKEDTNKWKNIP WGRINVKMAILPKVIYRFN PIKLPMTFFTDLEKTTLNFIVW KRARITKSILSQKNKAGGITLF FKLYYTATVTKTAWYWYQN MVLVPKQRYRSMEQNRALRN NAACLQLSDL WSPLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKBLEAFLTQKGKVGE KDDDFEKISELGAGNGGVFF VSHKPSGLVMARKLHELEIKP RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYCLLLPLD LCLSFSFSKVISDLVSDLHFGI KARNSHIGDVVVVELLPKNI WKGRTVALGENDC					l		HHDQVGFIPGMQGWFNIRKSIN
LKIIRÁIYDKPTANIILNGOKLI AFPIKTGTRQGCPLSPLVFNIV EVLAWAIRQEKEIKGJQLGKE VKLSLFADDMTVYLENPIISA MILKLISNFSKVSGYKINVOK. QAFLYTNNGQTESLIMSELPF ASKRIKYLGJQLTRDVKDLFK NYKPLLNEIKEDTTNKWKNIP WVGRINIVKMAILPKVIYRFN PIKLPMTFFTDLEKTILNFIVN KRARITKSILSQKNKAGGITLF FKLYYTATVTKTAWYWYON MVLYPKQRYRSMEONALRIK NAACLQLSDL H920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKGVKGE KDDDFEKISELGAGNGGVVFF VSHKPSGLVMARKLHELEE,P RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFSTSKVISDLVSDLHGI KARNSHIGDVVVVELLPKNI WKGRTVALCENDC			1				VIQHINRTKDKNHMIISIDAEKA
AFPLKTGTRQCCPLSPLVFNIT EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMTVYLENPIISA NLLKLISNFSKVSGYKINVQK: QAFLYTNNOGTESLIMSELPP ASKRIKYLGIQLTRDVKDLFK NYKPLLISIERDUTTKWKNIP WVGRINIVKMAILPKVIYRFN PIKLPMTFFTDLEKTTLINFIVN KRARITKSILSQKNKAGGITLE FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNRALRN NAACLQLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSIDI CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKGVXGE KDDDFEKISELGAGNGGVFF VSHKYSGLWAARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, RNOJIRSLGWARKLHELEIKP, KARRSHINGDVVVVSDLJHGI KARNSHINGDVVVVSDLJHGI KARNSHINGDVVVVSDLJHGI KARNSHINGDVVVVSDLJHGI KKRRNSHINGDVVVVSDLJKNI							FDKIQQPFMLKTLNKLGIHGTY
EVLAWAIRQEKEIKGIQLGKE VKLSLFADDMITVILEPRISAN NILKLISNFSKVSGYKINVQK: QAFLYTNNOGTESLIMSELPF ASKRIKYLGIQLTROVEDLFK NYKPILNEIKEDITNKWKNIP WVGRINIVKMAILPKUTYRFN PIKLPMTFFIDLEKTILNFIWN KRARITKSILSGKNKAGGITLF FKLYYTATVTKTAWYWYQN MVLVPKQRYRSMEONRALRS NAACLQLSDL MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKGKVGE KDDDFFKISFLGAGNGGVFF VSHKPSGLVMARKLHIELERP, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFSFSKVISDLVSDLHFGI KARNSHIGDVVVVELLPKNI WKGRTVALGENDC	1						LKIIRAIYDKPTANIILNGQKLK
VKLSLFADDMTVYLEIPRISA NLLKLISNFSKVSGYKINVQK: QAFLYTNNGQTESLIMSELPF ASKRIKYLGIQLTRDVKDLFK NYKPLLNEIREDTINKWKNIP WVGRINVKMAILPKUYVRFN PIKLPMTFFTDLEKTILNFIVN KRARITKSILSQKNKAGGITLE FKLYYTATVTKTAWYWYON MVLYPKQRYKMEQNALAR: NAACLOLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSIDI CSPLWNRTNLEALQKKLEELE DEQQRKRLEBALTQKGVCGE KDDDFEKISELGAGNGGVVFF VSHKPSGLVMARKLIHLEIKP, RNOJIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVISDLVSDLIHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC	İ		l	į	l		AFPLKTGTROGCPLSPLVFNIVL
VKLSLFADDMTVYLEIPRISA NLLKLISNFSKVSGYKINVQK: QAFLYTNNGQTESLIMSELPF ASKRIKYLGIQLTRDVKDLFK NYKPLLNEIREDTINKWKNIP WVGRINVKMAILPKUYVRFN PIKLPMTFFTDLEKTILNFIVN KRARITKSILSQKNKAGGITLE FKLYYTATVTKTAWYWYON MVLYPKQRYKMEQNALAR: NAACLOLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSIDI CSPLWNRTNLEALQKKLEELE DEQQRKRLEBALTQKGVCGE KDDDFEKISELGAGNGGVVFF VSHKPSGLVMARKLIHLEIKP, RNOJIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVISDLVSDLIHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC			l	İ			EVLAWAIROEKEIKGIOLGKEE
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QAFLYTNNOQTESLIMSELPF ASKRIKYLGIQLTRDVKDLFK NYKPLLNEIKEDITNKWKNIP WVGRINIVKMAILPKUYVRFN PIKLPMTFFTDLEKTILNFIW KRARITKSILSQKNKAGGITLF FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEONARLRS NAACLQLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSQDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKGKVGE KDDDFEKISELGAGNGGVFF VSHKPSGLVMARKLHELEKP, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFSTSKVISDLVSDLHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC			ı	1			
ASKRIKYLGIÓLTROVKOLTEK NYKPLLNEIKEÖLTROVKOLTEK NYKPLLNEIKEÖLTNKWKNIP WVGRNIVKMAILPKUYTRNN PIKLPMTFFTDLEKTTLNFIVM KRARITKSILSGNKNAGGITLEF FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNRALRN NAACLQLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGGGLYNSLGSI WSFLSMIGVLLWVDFSGDSIDI CSPLWNRTNLEALQKKLEELE DEGQARKLEEALTOKGKVGE KDDDFEKISELGAGNGGVVFF VSHKPSGLVMARKLHEIKP, RNOIRELGVLHECNSPYIVGE GAFYSDGESICMEHMVLLQE PCFGFLWMENSDYGLLLPNI LCLSFSTSKVISDLVSDILHGI KARNSHIRGDVVVSDILHGI KARNSHIRGDVVVSDILHGI KARNSHIRGDVVVSLLENNI WKGRTVALGENDC	1		1				
NYKPLLNEIKEDYTNKWKNIP WVGRINIVKMAILPKVIYRFN PIKLPMTFETDLEKTTLNFIVM KRARITKSILSOKNKAGGITLF FKLYYTATVTKTAWYWYON MVLVPKQRYRSMEQNRALRN NAACLQLSDL 14920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKOKVGE KDDDFEKISELGAGNGGVFF VSHKPSGLVMARKLHLEIERP, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFNFSKVISDLVSDILHGI KARNSHIGDVVVVELLPKNI WKGRTVALGENDC			l	1			
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KRARITKSILSQKNKAGGITLE FKLYYTATVKTAWYWYON MVLVPRQRYRSMEONRALRS NAACLQLSDL I4920 45288 A 15007 1 4215 MHSCLPDATGEGELYNSLGSI WSFLSMIGVLLWVDFSQDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKQKVGE KDDDFEKISELGAGNGGVVF VSHKPSGLVMARKLHELEE, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFNFSKVISDLVSDILHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC	ŀ						
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NAACLQISDL 14920 45288 A 15007 1 4215 MHSCLPDATGGELYNSLGSI WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEBATTOKQKVGE KDDDFEKISELGAGNGGVVF VSHKPSGLVMARKLIHLEIKP, RNOJIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLPLD LCLSFSFSKVISDLVSDJLHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC	ı						
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WSFLSMIGVLLWVDFSGDSID CSPLWNRTNLEALQKKLEELE DEQQRKRLEAFLTQKQKVGE KDDDFEKISFLGAGNGGVVFF VSHKPSGLVWARKLHIELEP, RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PCPGFLWMENSDYGLLLPLD LCLSFSFSKVLSDLVSDILHGI KARNRSHIGDVVVVELLPKNI WKGRTVALGENDC	14020	47200	 -	15007	1	4215	
CSPLWNRTNLEALQKKLEELE DEQQRRALEAFLTQKOKVGE KDDDFEKISELGAGNGVVFF VSHKPSGLVMARKLIHLEIKP, RNOIRELQVLHECNSPYIVGF GAFYSDGEISIGMEHMVLLGE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVLSDLVSDILHGI KARNSHIGDVVVYELLPKNI WKGRTVALCENDC	14920	45288	A	15007	,	4213	
DEQORRILEAFLTÓKÓKVGE KIDDFEKISELGAGNGGVVFI VSHKPSGLVMARKLIHLEIER, RNOHRELGVLHECNSPYIVGF GAFYSDGEISICMEHMVLLQE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVI.SDLVSDILHFGI KARNSHGDVVVELLPKNI WKGRTVALGENDC							
KDDDFEKISELGAGNGGVVF VSHKPSGLVMARKLIHLEIKP, RNQIIRELQVLHECNSPYIVGF GAFYSDGEISCMEHMVLLQE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVISDLVSDILHGI KARNRSHGDVVVVELLPKNI WKGRTVALGENDC			l				
VSHKPSGLVMARKLIHLEIKP, RNOIRELQVLHECNSPYIVGF GAFYSDGEISICMEHMVLLGE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVLSDLVSDILHGI KARNSHIGDVVYVELLPKNI WKGRTVALCENDC			l				
RNQIIRELQVLHECNSPYIVGF GAFYSDGESICMEHMVLLQE PPCGFLWMENSDYGLLLPLD LCLSFSFSKVI.SDLVSDILHGI KARNRSHGDVVVYELLPKNI WKGRTVALGENDC							
GAFYSDGEISICMEHMVLLQE PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVLSDLVSDILIHGI KARNRSHIGDVVVVELLPKNI WKGRTVALCENDC				I	l		
PCPGFLWMENSDYGLLLPLDI LCLSFSFSKVLSDLVSDILHGI KARNSHIGDVVYVELLPKNE WKGRTVALCENDC				I		İ	
LCLSFSFSKVI.SDLVSDILHGI KARNSHIGDVVVELLPKNE WKGRTVALGENDC				I	l	l	,
KARNRSIHGDVVVVELLPKNI WKGRTVALCENDC	1	1		ŀ	I		PCPGFLWMENSDYGLLLPLDLL
WKGRTVALCENDC					l		LCLSFSFSKVLSDLVSDILIHGM
							KARNRSIHGDVVVVELLPKNE
14921 45289 A 15008 I 2823	1						WKGRTVALCENDC
	14921	45289	Α	15008	1	2823	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
No.	sequence	liou	09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14922	45290	Α	15009	I	3253	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQGDLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
1						GSKALLSKWKRTEIITNYLSDH
						SAIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
1						RGKFIALNAHKRKQERSKIDTL
1						TSQLKELEKQEQTHSKASRRQE
İ						ITKIRAELKEIETQKTLQKINESR
1		l				SWFFERINKIDRPLARLIKKKRE
		<u> </u>				KNQIDTIKNDK
14923	45291	A	15010	I	2816	MEDEMNEMKREGKFREKRIKR
1		l				NEQSLQEIWDYVKRPNLHLIGV
1		1				PESDVENGTKLENTLQDIIQENF
		1				PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
		l				LRAAREKGHCNSGKSHRSLWE
						QNKGIFTTCICDDAEDLQSPSTA
						GPWFGVLYTKGGPYPQEDIFSE
						EVHTGPKLRKKIQEYQLTSKWS KSDVQVSVERRMAGGNPNQCH
		l				IGEVLLDGFTAEFYRRYKEELV
14924	45292	A	15011	1	3347	PFLLKLFQSIEKEG MGDFNIPLSTLDRSTRQKVNKD
14924	43232	l^	15011	l'	3347	TOELNSALHOADLIDSYRTLHP
						KSTEYTFFSAPHHTYSKIDHILG
						SKALLSKCKRTEIITNYLSDHSA
		l				IKLELRIKNLTQNCSTTWKLNN
						LLLNDYWVHNEMKAEIKMFFE
		1		1		TNENKDTTYHNLWDAFKAVCR
		l		i		GKFIALNAHKRKQERSKIDTLT
						SQLKELEKQEQTHSKASRRQEI
						TKIRAELKEIETQKTLQKINESR
		l				SWVFERINKIDRPLARLTKKKS
			ļ			EKNQIDAIKNDK
14925	45293	Α	15012	I	3855	MGKKQNRKTGNSKTQSASPPP
			İ			KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELWEDIQTKGKE
						VENFEKNLEECITRIINTEKCLK
						ELMELKTKARELHEECRSLRSR
						CDQLEER VSAMEDEMNEMKRE
	ļ.	1	1	I		GKFREKRIKRNEQSLQEIWDYV
						KRPNLRLIGVPESDVENGTKLE
1			1			NTLQDIIQENFPNLARQANVQI
			1	1		QEIQRMPQRYSSRRATPRHIIVR
		1	1	I		FTKVEMKEKMLRAARQKAPH
		_				HTYSKIDHIVGSKAL
14926	45294	A	15013	I	2559	
14927	45295	Α	15014	1	2956	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14928	45296	Α	15015	1	1853	MGRNOSRKAENSKNOSASSPP
						KDCSSSPAMEOSWTENDFDELT
						EVGFRRSVITNFSELKEHVETNL
						KEAKHLEKRLDKWLTRIKSVE
			l			KILNDLMELKTMAQELHDTCR
						SFNSQFNQVEERVSVIEDQINE
						MNIDGTYLKIIRAIYDK\PPRNII
				į.		LNOOKLEAFPLKTGTROGCPLS
						PLLFNIVLEVLARAIROEKEIKGI
1		l				OLGKEEVKLSLFADDMIVYLEN
						PIVSAQNLLKLISNFSKVSGYKI
				ŀ		NVOKSOAFLYTNNRQTESOIMS
						ELPFTIASKRIKYLGIQLTRDVK
l						DLFKENYKPLLKEIKEDTNKW
				ŀ		KNIPCSWVGRINIVKMAILPKVI
ľ						YRFNAIPIKLPMTFFTELEKTTL
				ŀ		KFIWNQKRARIAKSILSQKNKS
				l		GGITLPDFKLYYKATVTKTA/W
				ľ		L/YHTPLSCLPP*WNRTEPSEIIP
						HIMAVAQIQQVKLLLLQGQHA
				ŀ		SSRLQPTSSTKCGGWTASSASP
						RSMGLSSWKGNLASAIHYFPCK
l				1		PLTLPSRISTSSSSWGCRQAHAV
				1		LEVLRWPPGLGWPGWHHLTLL
Ì				ŀ		HVPPVRITKSNGPRACTRASAS
						HARHHQYNQEPSLSPLHSPLPP
]	l		PEQVLVSTAERTEDRSYHRTLC
						RHSPYFKNEAADLRGEGYSS
14929	45297	В	15016	1	3276	
14930	45298	Α	15017	1	2274	
14931	45299	В	15018	440	3383	
14932	45300	Α	15019	1	2646	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	Incation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
14000	145001	<u> </u>	1,5000		10000	MCKKONIDATONICKTOG + CDDD
14933	45301	Α	15020	1	2229	MGKKQNRKTGNSKTQSASPPP
İ			Ì			KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELREDIQTKGKE
						VQNFEKNLEECITRITNTEKCLK
						ELMELKTKARELREECRSLRSR
						CDQLEERVSAMEDEMNEMKRE
			ŀ			GKFREKRIKRNEQSLQEIWDYV
						KRPNLRLIGVPESDVENGTKLE
						NTLQDIIQENFPNLARQANVQI
ı						QEIQRTPQRYSSRRATPRHIIVR
		ŀ				FTKVEMKEKMLRAAREKEIQT
						TIREYYKHLYANKLENLEEMD
1						TFLDTYTLPRLNQEEVESLNRPI
1						TGSEIVAIINSLPTKKSPGPDGFT
			1			AEFYORYKEELVPFLLKLFOSIE
1			i			KEGILPNSFYEASIILIPKPGRDT
						TKKENFRPISLMNIDAKILNKIL
l l		1		1		ANRIQOHIKKLIHHDOVGFIPG
						MOGWFNIRKSINVIOHINRAKD
1		l		1		KNHMIISIDAEKAFDKIQQPFML
1						KTLNKLGIDGTYFKIIRAIYDKP
1		l				TANIILNGOKLEAFPLKTGTRO
1		1				GCPLSPLLFNIVLEVLARAIRQE
1						
		l	1			KEIKGIQLGKEEVKLSLFADDM
		1				IVYLENPIVSAQNLLKLISNFSK
		l				VSGYKINVQKSQAFLYTNNRQ
		1				TESQIMGELPFTIASKRIKYLGIQ
						LIRDVKDLFKENYKPLLKEIKE
		1				DTNKWKNIPCSWVGRINIVKM
		l				AILPK\IGKTTLKFIWNKKSRIAK
			i			SILSQKN/KAGGITLLTQLYYKA
						TVTKTAWYWYQNRDIDQWNR
14934	45302	A	15021	I	960	
14935	45303	A	15022	1	2928	MEDEMNEMKQEEKFREKRIKR
				ł	İ	NEQTLQEIWDYVKRPNLHLIGV
1		l				PESDGENGTKLENTLQDIIQENF
		1	Į.			PNLARQANVQIQEIQRMPQRYS
						SRRATPRHIIVRFTKVEMKEKM
		1				LRAAREKDRSMRQKVNKDTQE
						LNSALHQADLIDIYRTLHPKSTE
		1		1		YTFFSAPHHTYTKIDHILGSKAL
		1			l	LRKCKRTEIITNYLSDHSAIKLE
		1				LRIKNLTONRSTTWKLNNLLLN
		1	1	1		DYWVHNKMKAEIKMFFETNEN
						KDTTYQNLWDAF
14936	45304	A	15023	1	1782	
14937	45305	A	15024	ī	4881	
14938	45306	В	15025	I	5962	
		_				

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14939	45307	Α	15026	<u> </u>	2053	MKAEIKMFFETNKNKDTTYON
				ľ		LWDAFKAVCRGKFIALNAHKR
1			ł			KQERSKIDTLTSQLKELEKQEQ
1			İ			THSKASRROEITKIRAELKEIET
			l			QKTLQKINESRSWFFERINKIDR
						PLARLLKKKREKNQIDAIKNDK
						GDITTDPTEIQTTIREYYKHLYA
						NKLENLEEMDKFLDTYTLPRLN
						QEEVESLNRPITGAEIVAIINSLP
ł			l			TKK\SPGPDGFTAELYQRYKEG
			i	1		AVLEVLARAIRQEKEIKGIQLG
						KEEVKLSLFAGDMIVYLENPIV
						SAQNLLKLISNFSKVSGYKINV
-						QKSQAFLYTNNRQTESQIMSEL
1	ł					PFTIASKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
İ						PCSWVGRINIVKMAILPKVIYGF
1						NAIPIKLPMTFFTELEK\TT*KFI
1				ŀ		WNQKRARIAK*ILSQKNKAGGI
				İ		TLPDFKLYYKATVTKTAWYW
						YQNRDIDQWSRTEPSEIMPHIY
				ŀ		NYLIFDKPDKNKKWGKDSLFN
				ł	1	KWCRENWLAICRKLKLDPFLTP
				[YTKINSRWIKDLNVRPKTITTLE
				İ		EKLSNTIQDIGMGKDSMSKTPK
						AMATNVKIQKWDLIKLKSFCT
				1		AKETTIRVNRQPTTWEKIFATY
				ŀ		SSDKGLISRIYNELKQIYKKKTN
1				1		NPIKKWVKDMNRHFSKEDIYA
				1		AKKHMKKCSPSLAIREMQIKTT
<u></u>		_				MRYHLTPVRMAIIKKSGNNRC
14940	45308	Α	15027	3	3229	MACON TO LONG TO A MODE CALLED MA
14941	45309	A	15028	1	3907	MGDFNTPLSTLDRATRQKVNK
İ						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIITNYLSDHS
ı						AIKLELRIKNLTQNRSITWKLSN
						LLLNDYWVHNEMKADIKMFFE
						TNENKDTTYQNLWDTFKAVCR
						GKFLALNAHKKKQERSKIDTLT
1						SQLKELEKQEQTHSKASRRQEI
	l			l		TKIGAELKEIETQKTLQKINESR
1				l	1	SWFFERINKIDKPLARLIKKKRE
		<u>_</u>				KNQIDAIKNDK
14942	45310	A	15029	I	4005	
14943	45311	A	15030	287	2708	
14944	45312	A	15031	1	4602	
14945	45313	Α	15032	1	4806	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ 1D NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14946	45314	A	15033	965	5928	TWKGTTSTSRCKIMPKYRSTRQ KVNKDTQELNSALHQADLIDIY RTLHTKSTEV/TFFSAPHHTYSK IDHIVGSKALLSKCKRTEIITNY LSDHSAIKLELRIKNLNQSRSTT WKLNNLLLNDYWVHNEMKAE IKMFFETNENKDTTYONLWDA FKAVCRGKFAILAHRRKQERS KIDTLTSQLKELEKQEQTHSKA
						SRRQEITKIRAELKEIETQKTLQ KINESRSWFFERINKIDRPLARLI KKKREENOID
14947	45315	A	15034	383	8708	RECEEVOID
14947	45316	A	15035	2	3257	WRKIYOANGKOKKAGVAILV/S
						DKTDFKPTKIKRDKECHYIMVK GSIQQEELTILNIYAPNTGAPRFI KQVLSDLQRDLDSHTLIMGDFN TPLSTLDRLMRQKVNKDTGEL NSALHOVDLIDIYRTLHPSKTE YTFFSAPHHTYSKIDHILGSKAL LSKCKRTEIITNYLSDHSAIKLE LRIKNLTQSSSTTWKLNNILLN DYWVHNEMKAEIKMFFETNEN KNTTYQNLWDAFKAVCRGKFI ALNAYKRKQERS
14949	45317	A	15036		3508	MELKTKARELREECRSLRSPCN QLEERVSAMEDEMMEMKREG KFRÖKRIKRNEQSLOEIWDFVK RPNLRLIGVPESDGENGTKLEN TLQDIQENFPNLARQANIQIQEI QRTPQRYSSRRATPRHIIVRFTK KIDHILGSKVILSKCKRTEILTN TLSDHSAIKLERIKNLTQNHST TWKLNSLLLNDYWYNNEMKA EIKMFFETNENRDTTYQNLWD TYKAVCRGKFIALN
14950	45318	В	15037	1	4760	
14951	45319	Α	15038	1	3110	
14952	45320	A	15039		5546	MELKTKARELREECRSLSRSRN JEERVSAMEDEMNEMKREG KFREKRIKRNEQSLQEIWDYVK RPNLRLIGVPESDAENGTKLEN TLQDIQEOFPILARQANVQIQE IQRTPQRYSSRATPRHIVKPT KVEMKQKMRAAREKDFKPT KVEMKQKMRAAREKDFKPT KURNOKEGHYIMVKGSIQQEEL TILRIVAPNTGAPRFIKQVLSDL TURNIVAPNTGAPRFIKQVLSDL RSTRQKVNKDTQELNSALHQA DLIDIVRTLHPKSTE

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14953	45321	В	15040	11	4534	
14954	45321	В	15041	1	1826	
14955		_	15042	1	1626	
14955	45323 45324	B A	15042	248	1212	NEW LIVE CONTROL DEL DE LA LA LA LA LA LA LA LA LA LA LA LA LA
14956	45324	A.	15045	248	1212	NSKHYRSGQKKKLPTLPRAAN SRWRSLKI/GFTVSFDETLKPFV
						RDVSGSRLKDLPKPNKSDDETR
		1				
						ANDAVNRYKLLKKDARTIAAQ
						QVARLESAMCLRRRWSLENFQ
		1				LFLVEHPLVRHLTRRLIWGVYS
						AENQLLACFRVAEDNSSSTADD
l						DLFTLPEGDISIGTPHVLEISPTD
						AAALLFADYELLPPFRQLDRNS
		1				YALTEAERNASELTRWAGRKC
İ				1		PSGRVMGLANKGWIKGEPQDG
						GWIGWMIKPLGRWSLIMEIDEG
					i	FAVGMSPAELSAEQLLSKLWL
		ŀ				WEGKAERYGWGSNSTQEAQFS
		_				VIDAITASELINDIEALFE
14957	45325	A	15044	1	1788	
14958	45326	A	15045	1	3048	
14959	45327	Α	15046	1	3546	
14960	45328	Α	15047	969	1205	SPVRCCISAVHRTYHRRTSALL
		i i		1		RLAELHRLLEKCASRQWMYEN
ŀ		1				I*SPFRQIA*CLPYSDHKNQRLH
						GLDSIFLARLLW
14961	45329	A	15048	877	1248	WPLTVPYCGVGYAS*A*VADN
		ŀ				HRAVGYLSRWAGRKCPSGRV
		ŀ				MG/LANKGWIKGEPQDGGWI/G
						WMIKPLGRWSLIMEI/DEGFAV
İ						GMSPAELSAEQ/LLSKLWLWEG
		L.				KAERYG/WGSNSTQEAQFSVID
14962	45330	Α	15049	854	1313	DHGVASAATCVYTNEGHRQSY
						VRVCEYASRTRVSHHCSV*RVL
1		1				RCEAIRIVSPDRSC*RDPEFSYL\
		1				ALPENYNRLFLPNSTNQTNNRI
l						KTLNSIAIGKLLAAGGVYNGNI
1					1	EGFRDTAEKLDGRTIDGYDQIL
		_				NEKTAVIAKATA\SILLTKRS
14963	45331	Α	15050	1010	1396	KPFGKLTYKLFEGHAWLVKAE
1						KVRVGYRILGRLTYPFDIRFPAL
						PFGLVL*KGAHFTVPRIWAGHA
1			1			C*LMPMALPMFRWKVTGLLFIP
1						ICLVKPSFLTLTIIIAIRRISTSTN
	1					CRKTPKQPSRWCKPR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop cudun, /=possible nueleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequen ce	deletion, \=possible nucleotide insertion)
14964	45332	Α	15051	361	837	DGQQLIALHRLALRELQQAVH
		ŀ				AGLPQQAKILFDGGSEIGKIPSEI
i						YFCQLVKLILCPVLFPCWRRVA
						ILWRRRDVLFFGSFSFSALVSPH
		l				LCG/SYLPFVFDVGDRL*LSVG
		l				WS\KEIWPCK*GFGHLMSVLNT
						PMKSLVEGFKHFPLAGFGYEYL SFFCR
14965	45333	Ā	15052	1563	1652	STER
14966	45334	A	15053	2756	2995	AMRSAHRCTVFSALLNCWQITP
						HLTPSVMICGQLLTLVNRC*PSS
		l		ŀ		TIFSIIPLSKQVARMFRSAMSPL
		l		ŀ		NRARCWKVPCN
14967	45335	Α	15054	762	2184	WRAVPSEPISFILKADTEKYVY
		l				L*NSSSAQYSGSCTRAYYLDTQ
			İ			HPTPRLCLIF\GGKDSGLMLHLT
			}			AELARQMGKKICVLFIDWEAQ
			1			FSCTINYVQSLRELYTDVIEEFY
l						WDALPLTTQNSLSQYQPEWQC
		l l				WEPDVEWVRQPPQDAITDPDFF
		l				CFYQPGMTFEQFVREFAEWFSQ
l						KRPAAMMIGIRADESYNRFVAI
		1				ASLNKQRFADDKPWTTAAPGG
1						HSWYIYPIYDWKVADIWTWYA
		1				NHQSLCNPLYNLMYQAGVPLR
	i	1				HMRICEPFGPEQRQGLWLYHVI
		1				EPDRWAAIGSPADREEDAEEYL
		1				EAIMEARVTVAGMGLVMEVQ
		1				DYFDGEADRLAKAWLAEYTPQ
		1				IKSLKDERKEAYRQIVEMSTEP
		ı				QDVDLVRPANKFEMTRVREGE
						KEADLPVWKHHLLCDESGNYP
						ALLNHWETKVFEIETKREGFAF
		1				WYRNPQYTGQSSLGIAYVEAE
		<u> </u>				QYKIVRPDFLFFAEQDGKMVV
14968	45336	A	15055	2069	2586	PWLENLPVGICWSAVKKMGGT
		1				RSLVSCL/AVTLVSPQLMNAYL
1				1		LGQQLPEVWDFGMFSIAKVGY
	1	1	1	1		QAQVIPALYLVVVPVCSLILAV
		1		1		FLAHALIGPFGRMIGDGVAFAV
				l		RHLMTGSFAPIGAALFGFLYAP
l		1	1	I		LVITGVHQTTLAIDLQMIQSIGC
	1	1	1			TPVLPVNALSNIDLWHISILV_

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
i	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14969	45337	A	15056	356	1205	APASMTFEQFVREFA/VMVFTK
1						RPAAMMIGIRADESYTVLSPSP
l	İ					V*INNVLPTITWTFPQTIYDFLW
			ŀ			LIGEAIFF/YLPVGICWSAVKKM
						GGTRSLVSCL/AVTLVSPQL/MN
					ŀ	AYLLGQQLPEVWDFG\VCSLIL/
	i					AVFLAHALIGPFGRMI/GDGVA
						FAVRHLMTGSFA/PIGAALFGFL
						YAPLVIT/GVHQTTLAIDLLMM
						QSMGGTPVWPLIALSNIAQGSA
			ł			VIGHISSRKHNEREIPGEVFPRT
						GVGKGKSLG*PLH*LQPYRQHL
						GAGSYSQTLKKQIAESAVRLSIP
14970	45338	В	15057	1	435	
14971	45339	Α	15058	2	1785	WLLYRAHVKGE/STEALLPNM
	ł					VATSLAKLPIPKLMRWGATYV
	ŀ					HFVRPVHTVTLLLGDKVIPATIL
	ŀ					GIQSDRVIRGHRFMGEPEFTIDN
				1		ADQYPEILRDSGKVIADYEERK
						AKIKADAEEAARKIGGNADLSE
l						TLLEEVASLVEWPVVL/TAKF/E
l	1					EKF/LAISRTALVPYSADNMYQ
						LVNDVQSYPQFLPGCTGSRILES
						TPGQMTAAVDVSKAGISKTFTT
						RNQLTSNQSILMNLVDGPFKKL
						IGGWKFTPLSQEACRIEFHLDFE
						FTNKLIELAFGRVFKELAANMV
				İ		QAFTVRAKEVYSASMRYIYVIV
				l		YISYRPPNLLDSHIIGFSSITTHC
			l			AVTSHHYNSCDMTMTIHSYVT
						PYISVRVSLTQTYAPSPSRNLRI
		i		l		TDLTIYRAISSEKSIDDTRHIHTI
						ESRCRVCGAHDSRDAERLDRF
						AQTADEIRIADRGFSSRPECIRS
						LAFGEADYIVRVHWRGLRWLT
l		l				AEGMRFDMMGFLRGLDCALIS
l				1		KTRLLSENRRKGRVVQAETLE
l		l	1			AAGHVLLLTSLPEDEYSAEQVA
		İ				DCYRLRWQIELAFKRLKSLLHL
	1	l				DALRAKEPELAKVGIFANLLAA
		L				FLIDDIIQPSLDFPPRSAGSEKKN
14972	45340	Α	15059	I	993	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	,	
14973	45341	Α	15060	1	5621	MSEKTFLVEIGTEELPPKALRSL
						AESFAANFTAELDNAGLAHGT
						VQWFAAPRRLALKVANLAEAQ
		1				PDREIEKRGPAIAQAFDAEGKPS
		l		1		KAAEGWARGCGITVDQAERLT
		l				TDKGEWLLYRAHVKG\EGTEA
		l				LLPNMVASSVAKLPIPKLMRW
		l		1		GASDVHFVRPVHTVTLLLGDK
		l				VIPATILGIQSDRVIRGHRFMGE
		1				PEFTIDNADQYPEILRERGKVIA
		1			1	DYEERKAKIKADAEEAARKIGA
		1				KAWGTNVEFHVQNQP
14974	45342	Α	15061	82	732	
14975	45343	Α	15062	I	3849	MDLGANGWQTFRYVVLPNLSS
		ŀ				ALLAGGMLAFALSFDEIIVTTFT
						AEQVDAAVRAADAAFAEWGQ
			l	l		TTPKVRAECLLKLADVIEENGQ
		l				VFAELESSNNTAFARASSNGDL
						PTKADLQAQLDSLNKQKDLSA
		l				QDKLVQQDLTDTLATLDKIDRI
		l		l		KEETVQLRQKVAEAPEKMRQA
		l	ŀ			TAALTALSDVDNDEETRKILST
		1		l		LSLRQLETRVAQALDDLQNAQ
		ŀ	ŀ	ļ		NDLASYNSQLVSLQTQPERVQ
L						NAMYNASQQLQQIRSRLDG
14976	45344	A	15063	219	1186	PPKLGRKRGIFLPSHSNPNTNRI
						CPNCCMNGVSTANTTSKSSLVI
						CAWTKHALLNAILQPEQLWKS
			ŀ			GSRWRINLALPRGYRESLLQPW
					1	QAERLTTDKGEWLLY/RAHVK
						GESTEALLPNMV/ATSLAKLPIP
				l		KLMRWG/ASDVHFVRPVHTVT
						LL/LGDKVIPATILGIQSDRVIAA
				l		TGKAKIKADAEEAAR/KIGGNA
		l		l		DLSESLLEEV/ASLVEWPVVLN
		l		l		AKFDEK/FLAVPAEALVYTMKG
		l		1		DQ/KYFPVYANDGK/LLPNFI/F/
		1	l	l		VANIESKDP/Q/QIISGNEKSFVR
		l		l		LADAEFFF/NTT*KRLFLPHQKL
		1		l		PPSQILHQQTRNYASFCHRHRD
	<u> </u>		L	L		NWTSA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14977	45345	Α	15064	1	1496	SGGCAGLHGGLCSKEPFQASV
		1				QQGVAGHACAPAALQVPVCGT
						LSDCGSRAASLCENPSFISLYGR
		1				LVATSLAKLPIPKLMRWGASD
		l				VHFVRPVHTVTLL\LGDKVIPAT
						ILGIQSDRVIRGHRFMGEPEFTI
						DNADQYPEILRERGKVIADYEE
						RKAKIKADAEEAARKIGGNAD
						AKFEEKFLAVPAEALVYTMKG
						DQKYFPVYANDGKLLPNFIFVA
						NIESKDPQQI/DRIQALAGWIAE
						Q\IGADVNHATRAGLLSKCDLM
1		1	ĺ			TNMVFEFTDTQGVMGMHYAR
						HDGEAEDVAVALNEQYQPRFA
1		1				GDDLPSNPVACALAIADKMDT
l						LAGIFGIGQHPKGDKDPFALRR
			l			AALGVLRIIVEKNLNLDLQTLT
]		l	ĺ			EEAVRLYGDKLTNANVVDDVI
			ļ			DFMLGRFRAWYQDEGYTVDTI
		l				QAVLARRPTRPADFDDWHALG
ł		1				RAGTGDVVVGVRGRLAIVPLT
						DGLDEAGHSTGVLQGEVPQRL
						ERAATVPGVTLQPQALQARGL
14978	45346	A	15065	134	384	VTPTGWRKRRIRQWCSDAGCG
						VNALSGLRFSHSLALFSAAMFI
		1				LGVVSGITMSISP*ILSCVRSKNP
		_				LLVSGSGPLSCNSTIH
14979	45347	A	15066	1	723	TKSPKIARSRHIPLCHFIFERRPI
		1				MTNSNRIKLTWISFLSYALTGA
						LVIVPGMVMENIADYFNLPVSS
1						MSNTFTFLNAGILISIFLNAWLM
1		1				EIVPLKTQLRFGFLLMVLPVAG
ľ		l				LMFSHSLALFSAAMFILGVVSGI
		1		l		TMSIGTFLVTQMYEGRQRGYR
1		1		l		LFLAASLFSMAG\MIFPMIAAFL
				ł		LAPSS*W*WVYACLGLAYVTIF
1						IMTFGCEFPALGKHAPKSYAPV
		I.—			1015	EKVKWGILGGASSFRAV
14980	45348	Α	15067	422	1045	LLEGKLTNRKQ*HQH*QKARPC
						KKPIQSAQN/WNREEYDELTEV
				i		GRRWVIITNSFELKEHVLTQWK
1	1	l		I		EAKNLDKRLQEGLTRITSLEKKI
l		1		l		NDLMVLKNTARELHEAYTSINS
		l		l		RIDQAEERISEIEDQLNEIKLED
		l		I		KIRGKRIK/RQTNKQSL*KMWD
				İ		YVKRPNLRLIGVPESDEENVTE
						LENTLQDIIQENFPNLARQANIQ
		_				IQEIQTLLR
14981	45349	Α	15068	1	1131	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		-		sequence		
14982	45350	A	15069	383	756	LNPSNKLALINPAASPIIKGVDL
14702	13330	1	1.5005	100	/50	TANOI/PCIADIPLYDADVOOE*
ł	ŀ	l				CFPATVEALGEQIRQADGVVIV
		l				TPEYNYSVPGGLKNAIYWLSRL
		ı				RQILVFLDAMVMNKPEFMGGV
						IONKVDPOTGKVIDH
14983	45351	A	15070	343	924	IGHKYBIQIGKYIBII
14984	45352	A	15071	223	308	
14985	45353	A	15072	57	5546	NKMARKKFSGLEISLIVLFVIVT
		l				IIAIALIVVLATKTPAVDEISDST
		1				STPATTRVTTNPSDSGKCPNVL
						NDPVNVRINCIPEQFPTEGICAQ
!						RGCCWRPWNDSLIPWCFFVDN
1		l				HGYNVODMTTTSIGVEAKLNRI
1		ı				PSPTLFGNDINSVLFTTQNOTPN
		ı				RFRFKITDPNNRRYEVPHOYVK
		1				EFTGPTVSDTLYDVKVAQNPFS
		l			1	
						IQVIRKSNGKTLFDTSIGPLVYS
		l				DQYLQISARLPSDYIYGIGEQVH
14006	15251	_	15073	ļ,	1010	KRFRHDL
14986	45354	C A	15073	493	1818	
14987	45356	A	15074	381	498	
			15076	48	514	LLLFLLLQSYDWSESGKRQSAL
14989	45357	A	13076	48	314	
l		ı				HGRQKGLPKSPTAARGGSDPW
1		l				PSCRHP*GPNRKRRKS/R*PMET
1		1		1		APRRPSVSPALTPARPPGLTATP
1		ı			1	RLHPRPP/RSPNPSANLLRRPTP
1		l		i		DT*AHATFGPGRSRHHFPGQLS
		l				TE/RRRSAEGGRSRRNEASRRE
						GGE
14990 14991	45358 45359	B C	15077	249	1116 332	
14991	45360	A	15078	710	1326	D. I. O. IGEDEDOCTIVO C. I. I. I.
14992	45360	l ^A	15079	/10	1326	RAAGNCFPEPGFVKQLSAAAA
		l				VVAPAAAATEGSIPDTGKSPSSI
Į.		ł		l		SHWPNPQKNVICKRRGRVEQE
		l				QAWEKGR*RVVFGL*KSMRCP
	1	1				VDLLPQSCLCQRFLHQ/GKWIL
	1	1	l			TKDYIIHSAKSGRWLDATTYE
	1	1	I			WGYKIEKDSRYSPQMQSAPKR
	1	1	1			WREELKRTGAPGAFHRWKVVL
		1				LVRTDKRSDSLIRFSDNFGDCDI
						RMEEKTFRTLMES

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	locatinn of first codon for peptide sequence	endon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14993	45361	Α	15080	602	1257	FGKRGRTITLCILAVGMLCCRK GHTKPRRRPGASPISTHLHSSPPI GSCGAAAAR*PWWWNCDGWG PKCQRFSKH*DIAQPDEVSWGQ *PWAVGALAGPIPPPQASEGPL GCTPASICGSTRKPSKPI.SKKPP QAARTSSSRIVIFTPISTRPTTQ GPEKAKSATTNOLEGPSAKVK FVNFLMDETFPEPSFFI
14994	45362	С	15081	140	175	
14995	45363	A	15082	2295	3089	NLLGYQEINPPLQIFWOKLLAV SERDSSVKEESGDVESDDTML* MLPCP*ETTEFSS*PFRISS*LSS* STAQIPAETCFAGGWYSIFCPQ RIEQETMCLL*SWQSEVAVSCE SFWPTALHHSHFNSRAPPPCY/S KSTSSSLSSPLENSG*QNRPPSLL TISVLPLRITCLP*VSFIPIMQLG GSFLSNDRL*QSFLLPKNVSLEF VSFPLI*GAVSKPAISFNTSVSPL LVSDIFFALSLLP*GLGTWLIPSS SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLDTMSSLLST SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF SSLLLVRGF STERST STANDARD
14996	45364	В	15083	143	1727	
14997	45365	В	15084	307	1017	
14998	45366	A	15085	1	959	MSRMGILGFGERKELPNHPAV SEWPI-ILTVGSSGTVAEGEGN LTTSGLFSIVPLHPDGVAAGQR HSSLPRWGSQAEALLTSGTVDG QAEMLLFTDTMGGGEEMLLTS QTTGEDFITRSTESGTHSKSHKG DEGEPRWMLGWMLWVCDIAE PLVKAKLLDCKHTCCSVCJQ TRTSQKDVRCPWCSGVTKLPPD FSVSQLLDDPEVLAVTIPHASE TRCCPEI*AATCCPGTGRSPSLC DHPCWTAASARQGSPGGGGG GAQQAGHGEKLHVVEGVHCHL GGLRLGLPPQHRAP
14999	45367	В	15086	112	672	
15000	45368	А	15087	2	530	SGRSAALLSARVNRSQSFAGVL GSHERGPRSFPVFSPPGPPRKPP ALSRVSRMFSVAHPAAKVPQPE RLDL\VYTALKRGPDGLLGAVH QQEQEKLQGQIRESKRNS/RLG EWREGFLYDLDKQVKSIERFLR RLEFHASKIDELYE\AYCVQ\RR LR\DGALHMG\RAYTTGSPGSR

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15001	45369	A	15088	216	1369	EPVLRRPRQPRAGAQGFPVFS
						PPGPPRKPPALSRVSRMFSVAH
l			1	1		PAAKVPOPERLDLVYTALKRG
				1		L/QAYLEVHQQEQ/DKLQGQIR
			1	1		ESKRNSRLVMAGGVLWTVV*E
			1			RGAVTSLSASAGVLLIPPKYLPL
				1		LPTPAGLALYDLDKQVKSIERFL
						RRLEFHASKVRVQHVCRWD\D
						ELYEAYCVQRRLRDGAYNMV
l				l	l	RAYTTGSPGSREARDSLAEATR
			i	i		GHREYTE/DEGWGPMRQRHRS
				İ		MCLLES/ELEAQLGEFHLRMKG
			1			CAVDINDLGTIKLSLEVTGAAS
		1		l		SVNKASTVTKRSPPIARAHRTH
	1	1	1			PHFGNSFLSTRKPNTYTANHRR
		1				RPIEQHPDMRTPPPPHRARTNT
		1	1			GRKDKVRAHEGEKATPTRTRG
		1				APHGKAQRLRPLYQYVTIATLS
15002	45370	Α	15089	1	940	MSYNSKNYVETITTMNSVKFG
1		1				ETTPRNRTHYLSTNEKKIFLSGV
						FSNAYPAGLSRGWDSPWGSTP
		1				WVRHFCYLSSTQLCDQNCSLLS
		1				SLLPAGPSVCNRLLASAWQKSS
		1				SSASSEASETCQSVSECSSPTSD
		1			l	WSKVGSHEQPSGATLQRRKDR
ļ		1				VE\PCETQSPGPASGGTLGPSGE
		1				EAPRPRMSPATIAAKHGEEVSP
		1				AASDLAMVLT\RGMSL*HQKSS
	t	1		Ì		RDSLQYSSGYSTQTTTPSCSEDT
						IPSQGRPAGLGPTKPVSIESTYL
		1		į		QSPQRLKQEWATEGKIQKLPPQ
				Ì	l	IQVVNFQVDSSPSLPLIMVPEH
						MHVS
15003	45371	В	15090	1	777	
15004	45372	Α	15091	26	336	KISPMINVFCSESSPSGERRGA*
		1		l		TPPSPQSPPCGPKPSS\PPSCSPSR
1		1]		NYFLLFQGLCAEHPALAAPAAS
		1		i		VGTCPSPASGTHAAAPSGPWRG
	ļ	1_		l.,		RCVHSGPGGRLW
15005	45373	C	15092	46	423	DI DIVI IZZONICA DE CADO IL LIDO
15006	45374	A	15093	1773	2219	ELDYLYCSNSSAQCPGALLPSL
1		1		1	l	ALPGPAGSARQAGSPARRLCAP
l		1		I	1	VQPSGRWPAPGGPGFPGSSCPA
1		1		1	1	SPTRQ*LLTSGSGCPRPCHHPSG
	1	1				PPQ*QSTQAWDWVILGPGAHG
	1					PSGQRCERCLPSRPSGPQTHSG
1 - 0 0 -	45005	1-	15004	ļ.——	7016	VGSEPRAVRGPGVSFV
15007	45375	В	15094	1	7916	L

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hođ	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or pepude sequence	describin, (-possible intelecting ruser from)
ľ				Sequence		
15008	45376	Α	15095	16	545	GLPWRQHVRVEECGRS/PLRPC
						SPGTPSPS*PLPAATRGPATCOS
						LSGLOPEKTT/SPS/CAAVAHER
						PPGPVPQCPAQGTGLIF*EGWT
						OPOOIROGPHPVAHE\CAHDLO
			ŀ	ľ		QAAPGPGGAHQG/PSQPG*PDR
				ŀ		QPDAERAVVHPGAQ*DAEIQPG
						LPGEGGPLATRTERSPAORPSP
						OARG
15009	45377	A	15096	320	1167	MDLKTAVFNPARDGKLRLLTK
1.5005	10077	,	15070	120	,	*\MASKSKEEVSSLISEKTNGAT
						PLLMAARYGHLDMVEFLLEOC
1				ŀ		SASIEVGGSVNFDGETIEGAPPL
1				ŀ	!	WAASAAGHLKVVQSLLNHGAS
1						VNNTTLTNSTPLRAACFDGHLE
		1				IVKYLVEHKADLEVSNRHGHT
1						CLMISCYKGHKEIAQYLLE\KG
						ADVNRKSVKGNTALHDCAESG
						SLDIMKMLLMYCAKMEKDGY
				ļ		
		l				GMTPLLSASVTGHTNIVDFLTH
		1				HAQTSKTERINALELLGATLFT OKEISCDESGRTRDLSPRRKPPH
15010	45378		15097	187	623	
12010	45378	Α	15097	187	623	KLKTQKNNIGPQACNSQLASVS
						TRSPPPTLSVPIIMTDDSQASQL LLAAVTIRKLWSRMAPMLALP
						TGQSGQTGAEGQGKEGERGQS
		ŀ				PCWKPRLDPKLC/EPRQPAQLL*
	ŀ					ADVCTGP/CGVRGCPADCRLPC
16011	46270	<u> </u>	15098	789	1041	LSPSSAQQPKEASRSA
15011	45379 45380	A	15098	707	1373	SAWPSCHCCHLDILSR/PGDDA
13012	43380	^	13099	1,0,	1373	CSVOILVPGLKGDAGEKGDKG
	l			1		
	1					APGRPGRVGPTGEKGEKGDSG
	i					DIGPPGPNGEPGLPCECSQLRK
1						AIGEMDNQVSQLTSELKFIKNA VAGVRETESKIYLLVKEEKPYA
		l				
						DPPLSWQGRGGTLSMAQDEAV
						HGLKGAYLGQARLGRGFIGIHR
						PWKRRAPLLLF*TLSPLRSF*KG
						DGGGNTQKGSRKKRELAGQIW
		<u> </u>				GPCIRR
15013	45381	Α	15100	79	420	WRRLFLLGVLFRGYLGCLRSRS
	1			l		VLGRLKVSDMRIFFFACGCGHL
1	İ			l		STLPSWPLRPSLWLRL*EEQELP
1	1		1			SLSVPSSGRSASRLL*LGWRGF
						GPSPSPRLTVQFSGGRRAQGPS
15014	45382	Α	15101	3	284	
15015	45383	С	15102	83	112	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last endon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
15016	45384	Α	15103	84	303	EEQELPSLSVPSCLWGPCPRAG
						WLHSLHWPVSHSFSATSTFGM
						AIWT*SLASGIKPQSSSKVSSRIQ
						VPPTM
15017	45385	В	15104	112	462	
15018	45386	Α	15105	326	1323	DRKAINLQVKLEGAVAL*KP\G
						EAQLEFGKALGLFQ\STLGHHF
		1				VNNKEKILGDF\QEATLQAATK
		1				LTRGRSLVMENMEQLLSMRME
		1				DQSQRNMPLSVVLIREKAHSPF
		1				ENRKQEQGEGASPENFRAR\QG
		1				*FLAFKAAPPA*AASGPAAQER
l		1				SVOEPSVNAPG*IPEGGLHTLA
						GVHHE*ALAWCSPLNKPRFLW
Ì		1				KRLP/SRTFHFPVEDESAPGSKP
						SGDLLTLLLGGNAAGDFKLK\P
						CLVYPSENPCFLKGSFKPNLPL
						VWCSHKKAWVOPG*OSETPFO
						KKKKKKKKAWVTMSLFOEWF
		1				LHFCSTVE/ARCTQYDLPYKGL
						PILDSTPGHPGC
15019	45387	Α	15106	3	375	HASAHASGLQQSYQAVKEKSS
		1				EALEFMKRDLTEFTOVVQHDT
						ACTIAATASVVKEKLATEGSSG
					i	ATEKMKKGLSDFLGVISDTFAP
		1				SPDKT\STAMSSPYDGTKARLY
		1				SLOSDPATYCNEPDGPP
15020	45388	Α	15107	184	377	YCTNNQKVN*TKEIITKRKTPSR
						QKKDSAFSPSHPLTEGLKW\ND
1					1	LTLSVSLTHDGFVWRRGKGV
15021	45389	Α	15108	3	816	SWEDVGWWRSWLQQSYQAVK
						EKSSEALEFMKRDLTEFTQVVQ
l		1				HDTACTIAATASVVKEKLAIAA
l		1			l	CSRGACFLCPFSIQTEGSSGATE
l		1				KMKKGLSDFLGVISDTFAPSPD
						KTIDCDVITLMGTPSGPAEPYD
	1	1				GTKARLYSLQSDPATYCNEPDG
1		1		1	1	PPELFDA WLSQFCLEEKKGEISE
		1		1		LLVGSPSIRALYTKMVPAAVSH
1					1	SE/TFWHR\YFYKVHQLEQEQA
1		1				RRDALKORA/VSGDSCT*SGVL
	1	1				TNE*KGSWVRDTERVRSLPLQT
						FSEGVGWREC
15022	45390	Α	15109	1	281	
			4			

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15023	45391	A	15110		1194	MGFLHLGQAGLEFLTSGSSTDI SEDWEKDFDLDMTEBEVQMAL SKVDASGEPFCFWKGYPGGEL ETLIBLLQKLFFLKTDPAFVLSP STLVKLHILWGNEGLQESLESLV LHDAALVILVPNLEKKGEISELL VGSPSIRALYTKMVPAAVSHSE FWHRYFYKYHQLEQEQARRIA
						RLKQRAEQSISEEPGWEEEEEE LMGISPISPKEAKVPVAKISTFPE GRTLAPQSPCEENLVTSVEPPA EVTPSESSESIPKSPLPTRALGTD
15024	45392	A	15111	213	1031	AGNKPDPWAGRNRTAVLPDVS VFHREDVGWRSWLQQSYQA VKEKSSEALEFMKRDLTEFTQV VQHDTVCTIAATASVVKEKLA FGGSGGATERMKKGLSDFLGVI SDTFAPSPDKTI*\RDVITLMGTP SGTAEPYDGTKARLYSLQADP ATLLV*PUGPPELFDAWLSQFC LEEKKGEISELLVGSPSIRALYT KMYPAAVSISFEWIRYFYKVH QLEQEQAPRDALKQRAEQSIFE EFGWEEEEELLMGIPIFPKEAK FPLAKISYJP
15025	45393	С	15112	414	557	
15026	45394	Α	15113	98	467	
15027	45395	Α	15114	I	155	
15028	45396	A	15115	16	94	RCPIASEASWTITD*RCPIASEAS WTITDALGNSYSGGMAF
15029	45397	A	15116	95	115	IIRHLCND*TPREGCLPSP*PAW SDTFETWVNNQASLQ
15030	45398	Α	15117	3	282	LKPVITCLLQLGLLKPINSPYNS PTLPVQKPDRPYSKPNSLP*LNP SLLQRNYTSIFLLTLNMPSISCT TMLLYGQKEISSLHKGPPSLMP
15031	45399	A	15118	548	1401	SCTPCWHLTGRLWGITAVRQK RIVLSVDFKLRIQSRTGKAAFP WCLIIAGTPL*LYTHVSRVSDH AGMPALVLHPLQQVPLFWGRG NTL*KD*SLSLACVQHGLLKP INSPYNSPILPVVKPDKPYKLVQ NIKLINQIVLPHHVGGAPVLUQ TGGKSGKDCILWFECHLSHNTI EHQVSSSTFRIVVATGVLVSLH PQLYMAQNBERHAECLGESKG REGESLSGNPENSRFSLLTSRRY LYKSTRTIVLLGLAGPLKQKKL RSQHPSLYKYLQSFRNKDSYK

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15032	45400	Α	15119	144	298	WQVPLFWQPFPEPLELWPKA/H
		1				LTDSFPDLLGLAAED*RCPIASE
					\	APCTQVI
15033	45401	Α	15120	449	667	
15034	45402	Α	15121	2	111	
15035	45403	A	15122	323	401	LLLGLAAED*RCWIASEAL*TIT DAK
15036	45404	Α	15123	2	593	CTLPSH*NLITLTPLNANIPSHG
		1		l		ML*KD*SLLSLACYSTAF*GL*T
				l		PLTIPPFYLFKNQTRLTG*FRICP
		1				LSTKLFCLSTPWCQTHILSYPQY
						LPPOPIFLFWISNMLSLLFLCTL
		1				YPSLSSLSLGLTLTPIRLSKLPGP
						YCRKPSQTAPTSNHPSSFSGSW
	ŀ					YSVETSYPLSSSIFRKGRTDSWS
						LKDTPHQAQPPT
15037	45405	Α	15124	354	630	
15038	45406	Α	15125	29	354	RFLVSLMLPPAVVMLNHCYSLI
		1				FTNALVKRLFAWNMLSELATC
						AGNLATGPRNARSPGFLLSCVP
		l				SVRDPTGNQTVQLTWQPLPEPL
		1				ELWPKA/HLTDSFPDLLGLAAE
15039	45407	С	15126	1	576	
15040	45408	A	15127	377	624	RDQLQCLQLWDAGACYTCWK
		1				SATGPRNAHSTVFLLSHVPSVW
		1				DPTENRTVHLTWQPLPEPLELW
						PKA/HLTDSFPDLLGLAAED
15041	45409	С	15128	1	462	
15042	45410	Α	15129	44	340	PCQTTQGRLLTAGTPL*SFTHVS
		l				RVSDHAGTPALVLHP*RQVPLF
		l				WGRGNHISGTQELPNT*TAV/V
						QAF/LPEPPPTG/CLLHVPEIWPL
						GQGMPAGQDSS
15043	45411	Α	15130	242	522	RLVWPTATQRAWSQSEAAAAK
	1	I				TVDLALSARNSATGPRNACSPG
		ı				FLLSCAPSVWDFTGNRTVQLTR
		ı				QPLPEPLELWPKA/HLTDSFPDL
		1				LGLVTED
15044	45412	В	15131	1	1080	
15045	45413	Α	15132	1	642	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15046	45414	Α	15133	245	1033	TALLLTQSLFGGLFTQTRMKFG
						AVTRIGGPPLGDQSPVLLFFVPR
						ERSTYDLGPQTDQPKKHLTNFK
1		ŀ				STSFVFSSCIPPP*PSSISLLPPW/T
				ŀ	1	TDHAPLTISL/TT*SPLPCSMPIS/
						ASHSML*KD*SLLSLACYSMAF
	1					*SL*TLLTIPPFYLS*NQRRFTG*
	İ	l		ĺ		FRICALSTKLFCLSTPWCQTHIL
		l				SYPQYLPPQSIILFWISNVLSLLF
		l				LCTLHPSLSSLSLGLTLTPFRLS
1			Í	ļ	i	KLPGLYCRKASQTAPITSVKPK
						FLPYLLPISA
15047	45415	В	15134	146	556	
15048	45416	A	15135	1	555	LTSFLHCCLSHHYHGAQIHPRH
						FLHLICRCHCHQYCHHAYLDPC
	İ					PQDSHHLHWCWQPLPEPLELRP
						KA/HLTDSFPDLLGLAAED*HCP
						IASEAALPPCFPRPDMITSRLQL
		l				QRRRQTALGVRYRNATYRTRT
						ASTNSLRPRGSPFPLALVTNDR
		1		ì		QVRSQLALWKRDRLLQLSPINR
						AISESSFT
15049	45417	В	15136	1	555	
15050	45418	В	15137	499	769	
15051	45419	Α	15138	752	1111	MQPTIGNSQSVSRPTA*MPLAE
1						SHPITQLVGTVTTPRVTPLSIRH
1				ł	1	PWRCRSANAGAMILTMKQLAI
1		l				VMRVYQHQVRRSMRPGRNGA
		i i		l		RSGMSAVGAYYSSCIRVMTAL
		L_				QSCTGDDEAVLL
15052	45420	C	15139	1	632	DOUBLE CONTROL CONTROL
15053	45421	Α	15140	1	375	PGVRGAQGGPSIPRQCEESAIGP
]	1				KFAWIISSFHWKQGMFNMEAP
	1	l	1	l		SSLFFVNMCAVKKQTTWGRPD
		1				RGFIWQPLPEPLELWPKA/HLTD
		l		l		SFPDLLGLAAED*HCPIASEAPA
	L	ட	L		L	LVYTAGLHCFSKPSQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
15054	45422	Α	15141	1	1340	CSEYEDSSPAPVPATDLSSTLSS
						SVPQPQDTGTSQQLHPLDPWHE
				l		LLRAQELQGATNHKGYSHAEH
	1					EHAGLGVQGGNGALAFSNSGH
				l	'	RHAVPTISSGTGRRRTPSSSAFG
						LLNLHQWFVSGFQAFSDRLKA
				1		ALSASLLLRFGDSDWLPSSSAC
						KCLMLGLHFVIVGNICATLKEK
						YSSMLHLDVTMKKNGEKRTRL
	ŀ	ļ		ŀ		QKRKKGMPPHPAYEDLNIAAIT
						LPANVVLHQPSGFRTSGQLDPV
						WWSLDTDAHEIWCQDPGLGSG
						DFPWEITPLSSYSLLHEKDPPTT
						SGPQTDQPKKHLTNFKSKTKET
		ŀ				GFIHGPKTPAPVTDWEGSLPLV
						FNH\SGTPL*LFTHVSRVSDHAG
						MPALVLHP*RQVPLFWRRGKI*
		l				LTSPSRCTHEKSCNSLPPL*DKP
		1				QPHLQHTRTSKCLNRSGQAF/L
	l		1			PEPPPTG/CLLHVPEIWPLGOGM
						PAARDSS

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15055	45423	A	15142	2	2107	IDMIFTPGPPSTPKHKKSQKGSA
				l		FTFPSQQSPRNEPYVARPSTSEI
						EDQSMMGKFVKVERQVQDMG
				1		KKLDFLVDMHMQHMERLQVQ
				1		VTEYYPTKGTSSPAEAEKKEDN
						RYSDLKTIICNYSETGPPEPPYSF
						HQVTIDKVSPYGFFAHDPVNLP
	ŀ	1		i		RGGPSSGKVQATPPSSATTYVE
	1					RPTVLPILTLLDSRVSCHSQADL
						QGPYSDRISPRORRSITRDSDTP
						LSLMSVNHEELERSPSGFSISOD
						RDDYVFGPNGGSSWMREKRYL
				1	1	AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILQKQVG
		l	İ	1		QSITSMLGFLSRGPSMKLCMGL
1					l .	ACVLSLWNTVSGIKGEAKKEK
İ			ĺ			GMTFLPTTDSKKFFSLLSVTSYS
					i	SFAFHKFSVAVYNISNLKTVDP
	l					AKFPTRYCYCLNNRTNDLSDFT
					i	ALLVDIIGNSTSYLTEIFKSTSIL
İ		l				SVNOSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
		l		ŀ		SGVLALLLTQSLFGGLFTRTRM
1	İ	1		ŀ		KFGAVTRIGGPPLGNQSPSSCSL
		1			l .	LHEKDPPTTSGPQTDQPKKHLT
						NFKSAARPTFLGQGQVPLNPFS
		l l		l		FTLS/EQVLLS*AARTPOSLISTP
1		l l		l		QPLISVPQSLISVPQPLLYFSGG
		l l				QEPPPPPLLCVSSLFSRLASFTM
i i				l		
1	İ	l				GAFTHGTQTPSPTKATAPRYPQ TGDLSAEWPFTAGEEPVLVPRP
15056	45424	-	16142	,	020	TGDLSAE WPF TAGEEP VL VPRP
15056	45424	В	15143	401	930	
15057	45425					
15058	45426	С	15145	1	843 498	
15059	45427	A	15146			CLOCKLED TRANSPORTER
15060	45428	Α	15147	112	669	SLQGRLSDYTPTFQGCQTTQGC
	1	1				LPWSFTLCSKSRFSGEGETRFKR
	1	l l				IKACYHSPATAWPFKAYKLSLQ
	1	l				FPHFTCRKTRQALQVSSESAPY
		1				QPNCFAYPPRGAEPIHSPILNTS
1		1				LHNPLFCSGSQTCFLY/SFLCTL
1		1		l		HPSLSSLSLGLTLTPIRLSKLPGL
1						YCLKSSQTAPITSVKPKFLPHLL
				l		PISA
15061	45429	Α	15148	957	1100	

SEQ ID NO:			SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	noa	09/540.217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	, , , , , , , , , , , , , , , , , , , ,
15062	45430	A	15149	303	2028	LGSGDLPWEINPLSSCSLLREKD
13002	45450	<u> </u> ^	13143	1303	2028	PPTTSGPQTHQPKEHLTNFKSD
		l		l		KGDMFYPWTQN/YQCRSRTGK
ŀ						AAFPWCLITAGTPL*LYTHVSR
ŀ						VSDHAGMPALVLHP*RQVPLF
						WGRGKLLLARPS*VPILPOPLLL
						HPIIFLSPPLLTPGPAYSFVL*LA
						LPHLPSNLLLKRWLEPKA*SSM/
						TLKGLKPVITRLLQHGLLKPTN
						SPYNSPILPVLKPDKPYKLVOD
						LRLINQIVLPIHPVLGIIGLTSSV
1						RRDAGODLKRDRAEFLLGDEV
						HHPHRRRIAEARRLLLGQHFDP
						LHRLIGQVLELGEARYAPPVEQ
					1	HHRLAPARRTGQRLHPLEQFG
						QAGRAQRRNRLGIEHRDRLDRP
						DDGAGNALAGDGDFRRGCLFA
						GIGIRPRYCRYRQSQDDRRPSH
					İ	AAPRPRALPIPHRPATPIADVM
1		İ				VMFSVAIMSQLRLVLRHIDGIS
						RWPPAVVFITDVKIWLVYPSDD
1					l	CRIRSNDRDDMQGEAPAMSMN
						AAARVGDPIGHSFSQGLFGEAL
1						DGLFFARRSEVDMRAGNLGRLI
					ŀ	ARGLSGGRWTPADGQLTLGSR
						DVFINGPPATMTIRSTGQCRQH
15063	45431	A	15150	653	1076	RSKYPNLVSLCPSPLFPRPDLLS
						LWPNPLFLHPNLLYLCAPIPYFH
						APTSYLCTPTPYFHDPTPFPLFW
						KELATSARNLATRPRNACSPGF
						LLSCVPSVRDPTGNRTVQLTW
						QPLPEPLELWPKA/HLTDSFPDL
						LGLAAED*CCPIASESP*TSDILG
						RDTLLLALKVQTVVLQTACGE
			1			GHVAGNCGRPLETEGSLQLTAT
			İ			KKLRDSVLQPKSPEFCQQFTRA
1	ļ.					WNRTQVPDETEAPAGTYAAQS
1						GDLPWEINPLSSCSLLHEKDPPT
1						ASGPQTDQPKEHLTNFKSGFRG
						VRPRRDACLGPSPLAASPAFLG
						KGQVPQPLLSVSLPLLRLSGGQ
1	l		l	l		ETPNPFSFTLSGKSAFLEEQVPQ
	ľ		1			PRISVPQSLISTPRPLISVAQSLIS
	l .			i		APQPLISLCPNPLFPCPDLLSLHP
	[l		NPLFPRPNPFPAFLEGACYKCQ
						KSGHQAKECLQPGIPPKLRPICA
						GPHWKSDCPTHLAATPRAPGT
15064	45422	C	16161	234	449	LAQGSLTDSFPDLLGLAAED
15064	45432	U	15151	234	449	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15065	45433	Α	15152	2095	2663	LGLGDLPWEINPPSSCSLLREKD PPTTTSGPQTYQPKKHLTNFKSA CFERIKACYHSPAKAWPFKAYK LSLQFPHFTCLKTRHAI/PS*FSD LCLINQIVLPIHPMVPNPYTLLC SIPPYTMHYSVLQLIRAFFTIPLH
						P*\PSLY*VSLGLTLTPIRLSKLP GLHCRKASQTTPITSVKPKFHP HLLPIST
15066	45434	А	15153	678	1189	LGSGDLPWEINPLSSCSLLCEKH PPTTSGPQTDQPKKHLTNFKSE TKETCFIHEPKTLAPVTD*GRQP SLGV*SLQGRPL*SFSHISRVSD HAGMPALVLHP*RQVPLFWGR GKYPNPFSPCLLPLLCFSGDRG KYPNPFSPCLLYSFSAFLGTGQVP QPLLSLSLPILLCF
15067	45435	А	15154	649	1428	GCISELOFLLASTVRQTPATSPA HKNFQTPELQRPGVPPEPPPTG ACVTCRKSGHWAKECPQPGIPP KAPHLTDSFPDLLSLAALED*HC PHASEAPETIDAELEVTLTVEA HLHPGEINSHVAHTKPVWWSL HRDAHEIWCCDSDRGTSLRRPI PCPALCSVTKHILRPQVLRFI PRNISPISNPAAPHQAELGPNPS/ SSLCSSTL*SFYHLPSSHLVWLT VSFRD
15068	45436	A	15155	240	1208	LRSGDLPWEINPLSSCSLLREKD PPTTSGPOTHQPKEHLTNFKSC KQ;DLFLLSSPTSLTIPQPLSPFN LGATLQSLPSLNFNSFHFLVETK ETREFIRGPKTPAPVT;VLGROPS LGV*SLQRCLSDYTPTFQGQT TQGRLPWSFTLSGK;CPLFWGR GNTL*KO*SLSLACYSMGLLN PINSPFPSSHFTCPKTRQAVTS* VQDLRLINHIVLLPHIPGGAQPH TLFCPQVRPVLPHILFVYVKHA FFTIPLQPLIPSLSLPSLRLTLTPI RLNKLPGLCCCKAFRDSPHVLN QAQISSSSVTYLGIILIKAHVLSL PIVSDW
15069	45437	Α	15156	1174	1623	KFGLVQLTLGKPLPEPLELWPK A/HLTDSFPDLLGLAAED*HCPI ASEAPYTNTDAELRVTLTVEGK SVPFLINTEATHSTLPSFQGPLSL ASITVVGIDGQASKPLKTPQLW CQLRQYSFKHSFLVIPTCPVPLL G*DTLTKLSASLTIP

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
		<u> </u>			liero .	Mar avar av un illi il name
15070	45438	Α	15157	1	1610	MGLGKSLCILKDAVLALREIKG
						ISFQYAPNTNRLWGYRQAKQT
						QHLSGISTLDCPQRYNGDDTGS
				l		VWALEPSCMLWACPLVFPSEIY
						YFISEDVWKHLDAQEKFAAVM
						GPSRRTSVRAVWKGNVGLEPP
					ľ	HRVPTGALPRVAVKRGPLSSRP
						QNGRSTYGLHCLKDTPTQEDW
	ŀ					LVSVLPEGSRVGVDPLIIPTALG
						HGRTTFLKVLDFQGPAPCPAA
						MGVRLMAGPQCVSSDYWKKM
						AKVLRSAGHHLIPVKENLVDKI
			i			WTDRPERPCKPLLTLGLDYTGI
						SW\KDKVAD\LRLKM\AERNVM
			ŀ			W\FV\VTALDEIAWLFN\LRGSD
						VEHNPVFFSYAIIRD*ETFMLF\I
						DGD\RIDAPSV\KEHLLLDLGLE
				1		AEYRIQVHPYKSILSELKALCA
				1		DLSPREKVWVSDKASYA/V*SE
				ľ		TIPKDHRCCMPYTPICIAKA\VK
						NSA\ESEGMRISDEKSAGNLTED
1						PFYVIDRTLITTGVVCNSTEKSS
						LMSVASADTAENLWHLFFHILP
						SERALTRKLRVLKLGPVNYSVO
1						NPTAQLHQQGPHLGVA
15071	45439	A	15158	1	756	
15072	45440	Α	15159	1	1677	
15073	45441	A	15160	1220	2620	GPTEPHTGELWLESGGCPSGTK
1		1				LPEEGTGSNLCCSTASAGDTQA
						NRVCSGPPTNHSRPAADRPVRR
						KINKOKTATSTSTKRTSTORPH
						PKVTNDHSSSPAREONWMEN\E
						FDELTEVGFRRWVITNSSELKK
1						HVLTQCKEAKNLEKRHSLGTK
						EPAALKGRTOYWODSSPADLR
						PWLCATLLKQYKPTVFSHSTVA
						VHPRAGFLNLSTVDOHPVLDEP
						IAEAVCIIADTNKWSVQVATSQ
						RKVPHNLKLGQDVLVSSQVSSL
						LOSILOLYKLHLPADFVGDILLD
						NNLKCVFOLGSILPVTFRSHDP
						YDGEGTDELRTAACSPDTEQLA
						RQSGTYPVSLGQATVLEDSKA
1	l	1				
	l					ALPGLGPTLMAVTVVTTEAAV
						DMVVGDRGEERGGTLVAPGTR
	l					GLTEEIWFPVPSPSRWWSSHPV
	l					VAFITLAKYGPTSSPPVFHHGPN
						IIRHGAQGAASAGDSGSSSRCD
L	L					RLPCRQNA

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence	1100	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		037510,211	sequence	or pepade sequence	
ł						
15074	45442	Α	15161	666	1536	FSSAGDTQANRVCSVPPTNHSR
						PAADRPVRRKVNKQKTATSTS
						TKRTSTORPHPKVTNDHSSSPA
		ŀ				REONWMEN\EFDELTEVGFRR
		l				WVITNSSELKKHVLTOCKEAK
		i i				NLEKRVRVNCLT\RIT*P*RRTI\
1		ŀ				KTLMELKNTARE\LHEAYTSINS
1		l l		1		WINOAEERLSEIEDOLNEIKRED
						KITEKMKNSTTVRVAASMOSK
		ı		1		LLOGVAEEGPLRLTRSASFSAEI
		1				SMCHQWLFSTFGNGKSRNRYH
		1				CEKAHIEGNQLVEETGRAFKSR
		l				HRYGTGPLHKFEIHEYTELHHV
1		l				
		_	1 4 4 4 4		261	LMLQGKC
15075	45443	С	15162	104	364	
15076	45444	Α	15163	162	467	
15077	45445	В	15164	1	1122	
15078	45446	A	15165	188	456	CECUTA DI LI LETTE CONCOCO
15079	45447	Α	15166	485	1580	GEGYKADLAAATVECPICQQQ
1		1				RLTLSPQYSHIPQGDQPTTW*Q
i		ı			ł	VDCIGPLLKGQRFVLTGIDTYS
1		1				GYVFAYPVCNASAKTIISGLTE
1		l		i	1	CLIHCHGIPHSIASDQGTHFMA
1		l				KEVWQWAHVHGIH*SYHVAD
1		1				YPEAAGLIEGWNGLLKSQLQC
1			ļ			QLGDNTLQGWGKVLQKPMYA
1		1		1		LNQHLIYGTVSPIATIHGSRNQD
1		1		1		WNGSSLRGWTFTILFTAPRREP
1					i	GTYIASDQRTHFMAKEVWQW
ı		1		ĺ		AHAHGIHWSYHVADYPEAAGL
1		Į.		1		IEGWNGLLKSQLQCQLGDNTL
1			1	l		OGWGKVLOKPMYALNOHLIY
1		ĺ				GTVSPIATIHGSRNQGMEVEVA
1		1	İ	i		PLTITPGDPQAKFLLPVPMTLCS
1		l				AGLEVLVPEGGTLPPGHTTIPLN
15080	45448	В	15167	10	2352	
15081	45449	A	15168	2	497	WKPIAMRKSRRRQEKLLPVQK
						EGOPKEGEREEGGPVOAOKVP
		l		į.		NRLHLHWTT/HLPSFFOLPLLLD
		i				SLWRALKFVVLEAIVKMIRFNG
1		I	1	l		VLGKCEAIRAOALHEGGPPLPL
1		l		l		AVFHNRRLQIVACSRISARLRGS
	1	l		1		
				1		PPPKSNNEQSSLSTDLRTDSQRL
15000	45450	<u>_</u>	15160		1011	PPLSRNAQQKG
15082	45450	В	15169	1	1911	

SEO ID	SEO ID NO:	Mct	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
			1	sequence		
15083	45451	Α	15170	1	1569	MPIGKVGEYYIKGTLRATKESE
						QQSSALELSSDRVYLNEKEPGD
		1				QPWNAAFVFRGCGVALALRPP
		1				HLGWGEPRWAGFRGAFGFPFS
						TLNPSNWSLTLIIQGCKQEGFYS
						PSMRPRVVPKRGSAPKVRAEEN
İ			1			AALPSRCPPGPLPVAQPRGLLM
l		ŀ				GAHVSSAPAVGVGAASCHRLC
						PGFPDRPRPTLRSCIMNAVVGG
İ						CDLRGQVYDPTGGWANEGTR
l						DLSAVQGVPNHPGGRPCPTPQS
ı						SRQLPVVSRRPVVRPGGEAPPP
	i		l	ľ		LDILPAALALHQSLLTPQDPRE
		l	1			DRPSPADLWAQDSWVGQPQEL
						PVGSRVRTGTEPDTHSGSVCTS
l		1				RQGKDGSGPLARESSVRVILQG
		1		1		PSSAWMMFPEYTPGPRARAGT
l		ĺ				AVVGAMARQTVNVCLTKVRL
ĺ						QHPEGIPGGGRGPLVPKPVCTC/
						IVDVKGSRV\WHVHICMSMGC
				1		RHVCLCGRSVGSWRMCTAGQT
				1		PKLARQASIELPSMAASSTKSW
						WETGEVQAQSAAKTPSCKDIV
						AGDMSKKSLWEQKGGSKTSST
		١.				IKMDSVTT
15084	45452	Α	15171	328	641	LLMRSWSLPSAPRGFSCSCLQP
				l		QYPCRTESWPQTEPVPSPVSHR
				İ		LLVLATAMLGSSME/RLAG*LG
ĺ				ŀ		GPASGLDGLGVLFQPLINLGDG
ĺ						QVWLGFLHTVTLLYGAV
15085	45453	В	15172	120	1458	
15086	45454	Α	15173	1	440	IQQDSQAPLYS\IKETQRANTYL
			1			VEWEPGAETAFKTLKQALVQA
				1		PALSLPTGQNFSLYITERAGIAF
			l			GVLTQTRGTTPQPVVYLKDGEP
						IEHDYQQIIAQTYATQDDLLEV
						PLANPDLNLYTDGSSFVVNGIR
		L				RAGYAIVSNVTVLE
15087	45455	Α	15174	288	376	NHWLLLTMDPRIQQDSQAPLŸ
1		1	1			S\IKETQRA

SEQ ID			SEQ ID NO: in USSN		Nucleotide location of last endon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	09/540,217	location of first codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		0,7,0,40,217	sequence	or pepinac sequence	
15088	45456	Α	15175	777	1548	SKIQRDSQAPLYS\IEETQRANT
i						HLVEGEPEAEIAFKTLKQALVQ
i						APALSLPTGQNFFLYITERAGIA
						LGILTQTRETTPQPVAYLSKEV
						DVVAKVWLHCLRDGEPIKHDC
				ŀ		QQIIVQIYAARDDLLEVPLANP
		l				DLNLYIDGSSFVENGIRRAGYAI
						VSDVTVLENSLAAVTLQNRQG
i						LDLLTAEKGGLCTFSGKECCFY
						TNQSGISSPLEDTTTAGPFLHPI
		l				QQEVARAIIGQFPTAFGVSCLE
		i		ł		GRLRGEASWTSWVE
15089	45457	Α	15176	1	1359	MGPEPKNTHPGSCTCPLACSSS
1		1				HKGFECLAVEQGSYTPVASPIV
1		1				GVKENQEILDLLELKYNQTRSA
		1		l		VTRCVLGAASTARIKQAEEGGR
		l				SRLAESSGLRLPFVLELPALEHQ
1				l		TPSSSGFELLDLHQRLATWSQA
1		l			l .	FNYRLKDTLLASLLLRFWDSD
		ı				WLPCSSVCSWPIVGLHLVIIERN
		l			1	YSYQQGTETVVSSWATGVSAA
		l			1	ASVSATIPINRVQGHCGFLGKR
		l				CFYCCIGECNHSDHQGPGTVAG
	1	1		1		SWARGVSATASQIQVDLGKFS
		l			ł	DDPDRYIDVLHLTWRDVMLLL
		1			ŀ	DQTLAFNENNVALAAAQEFGD
		1				TWYHRPVNDRMTAEQRDKFPT
		l				DIKTAAGIPWNHWLLLTMDPQ
!						TQRDSQAPLYS\I*ETQRANTYL
		l				VEWEPEAE/TSFKTLKQALVQA
		l				PALSILTGQNFSLYVTERARLAL
		l				GVLTQTCGTTPLYQDELRKKTL
		l			i	RHRDSEGSGFNQLEASAD
15090	45458	Α	15177	368	5192	AAPAPQLGLGPCKLASHLLSQS
i		l	ŀ			HLSQSVRSEAEQKAGASPASSI
		1				QREPLFKDQLKAPLPEQKRRSL
	1	l				LTVPLLTFRAQKCGPGSPLVAR
	1					GGATDSWLVSEEFRLSDSTAPP
1	1	1	l	1		TELRGQEATGITKPPASLHPQPH
		l			1	AWDWARGRDPRRHLLGTPELQ
		l			l	EHSEKTTTETLPADMGKLRTW
		1			ŀ	ERKEATAGQAAEGAAGGIHFE
1	1		l	1		ATSLLRPSSYVPQTQGALSTTY
	1	1	l	1		KEAKMDRDLHLTPILAAQRAR
	1		l	1		GPRIHSFPSDEMIEPS
L		_		L		

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15091	45459	A	15178	76	969	KKSTALVIDNGSCMCKAGFAG DDAPQAMFPSTVGFPGHOGVM VDMGQKDSVYGNEAQSKRSIL TLKYPIKHGIITNWDNMEKIWH HFYNELHGAPE*QPVLLTEDPL NPEDNREKMTQIMEETENTPA MYVAIRDVLSLYAFGRTTGIVM
						DCGHRVTHTVPIVEGYSLPHAI LHLDLTGWDLTEYPCILLFSEK SYELPDGQVITIGNEWFWCPEV LFQPSFPCMESCRIHQTTINSIKK CDVDIRKDINTVLSGGTTMYPG LDAPMGKEITALAPSTMKIKIIA PPERKFSVWIGCS
15092	45460	A	15179	1	451	
15093	45461	A	15180	130	227	LMKMKKVM*RA*KEEERRTPP AEDDFEKDLEWLINENEKSDAS IIEVYIANSIHNNYITKCFAIL
15094	45462	A	15181	104	1715	IEEERRTPPAISVIPAARPYTAY FLHKQKMNENKOTDSKKSEEY EDDFEKDLEWLINENEKSDASII EMACEKEENINQDLKENETVM EHTKRHSDPOKSLQDEVSPRRN DIISVPGIQPLDPISDSDSSNSTG ESKLESQKDLEEEDEEVRRYI MEKIVQANKLLQNQEPVNDKR ERLLKFKDQLVDLEVPPLEDIT TSKNYFENERNMFGKLSQLCIS NDEGGEDVLLSLTNGSCEENKD RTILVERDGFELINLODIASQ GFLPPINNANSTENDPQQLLPRS SNSSVSGTKKEDSTELLNLODIASQ GFLPPINNANSTENDPQQLLPRS SNSSVSGTKKEDSTAKIHAVTH STGEPLAYHQOPPLNRKTCPSS AVNSDSKGNGKSNHRTQSAH ENDIVFKAWLQKFLEERKKR ENDIVFKAWLQKKREQVLEMR RIQRAKEIEDMNSRENRDPQQA FRLWLKKKHEEDMNSRENRDPQQA FRLWLKKKHEEGMRERGTEEL WLRKKKHEEGMRERGTEEL WLRKKKHEEGMERGFREFAFKQ WLRRKEMEKMAEQQAVRERT RQLKLEAKRSKQLQHHLYMSE AKPERTIDHYN

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
	l			sequence		
44004	10110	<u> </u>	1.5100	I	2040	My Lournomous vo Look van
15095	45463	Α	15182	881	2040	MKAGWRQTQHLHDAGQLLHL
						LPGEQGVARVQLRRDAAQAPH
						VNGHVVGVAQDHFWGSVEPA
	ł					LDVRVYCGGKTGGKQRERREG
	l					RCGVEILMVGERKMNENKETD
						SKESEEYKDESEKDLEGLTNEN
	1					EKSDASIIEMACEKEENINQDLK
					l	ENETVIEHSKQLSDPDKSLQDE
		l			İ	VSPRRNDFISVPSIQPLDPISDSD
		1	1		l	SENSFQESKLESQK DLEEEEDER
	ŀ	1				VLPRSSNSCVSGIKKEDSAAKID
		1			ŀ	VVTHSSTGEPLAYI/ASTYKLAY
		1				IPPLNRKTCPSSAAISDRSKGNG
İ	ł					KSN/HGQSAHISSVTSTYCLSP*
	1					QKEL/QKREQKREKLKREEEQQ
	}					KIEEEEEKKRENGIVFKAWLQK
1						KREQVLEMRRIQRAKEIEDMNS
		L_				RVSKTSLKNKFIQI
15096	45464	Α	15183	873	4292	MVVSLGRVSSLSEDSDPTGRVG
ŀ	1				1	R*VDCK VDRSIWAGVWISEDAF
1	ŀ					SEA*EIGGLGRK*MLSGGPQSPS
	ł	1				KHVHIL*DQDSSSIPEPMGPCCP
	}					PHAPGKEPGTSCSIDTDGKLSG
	Ì			l		VSGDDS*T/GTDGGESEEILLPQ
						*QLPVTLPSLPLPG*VRYTSRSR
						SRSRSTRDPGRKSRPISGLRGLC
		1		i		GDGATS/PLTPRHAPARDVQ/PG
		1		1		VSMAFAGDR\GDGGRRGGASQA
		l		ı		PAPPGPAVPRPCAQYSARAMPA
	10160	_	1.004		2214	SGRRES
15097	45465 45466	B A	15184	3	2214 361	
15098	45467	A	15186	1	1035	MKIHKSGGLKATPQLYASQFSG
13099	43407	^	13100	ľ	1033	VAILATWGPHRPRNSSTSFSTOS
i						KLRRPDESSVDFOKLMLCKEAT
						KKSKEKEPGMAFPQGHLIFSDV
		1	İ			AIKFSVEEWKCLNPEORALYRE
		ł				MMLENYRNLES/VG*LFKIHDG
		1				VLINRARQYRSVPHRDIGKT*FL
		İ				LPRNQERYSGL*VSVARN*KK
						WP*STHDRNORVN*VHTGEKP
						OKCNECGKTFSOKSYLOCHHR
1	1	1]			LHTGQKPYK*E*CDKVYSCRSQ
						LKTHRRIHTG\EKAYKCKVÇEK
						AFWDNSCLSCQKRVHIGEKPYT
		1		I		ALVMHKAIHTGEKPYTCNECG
				i		KAFSRKANLALYHRLHTGEKP YKCEECDKVYSRRSHLERHK
15100	45468	В	15187	1	1347	I NUEECUK V Y SKKSHLEKHK
12100	42408	٥_	12197	11	1341	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i		sequence		
15101	45469	A	15188	l	1446	MWRETKRYAAVLLLTPHYSAS
13101	43409	<u>۱</u> ^	13100	1	1440	LLPDW/PNLYLHQSGHHVPGVY
						LKFG\SSNPIYPCVDSAKTVFPN
						APLHDRVOSVRRMGPFKGVFR
						KVFSNLGFEYSLSTWATMGSPG
						SLGVSPFGKTLFVESASGSLDA
						LWPTVEKEISSHKNYOKHSEKO
		l				IYDVRIELTGLNALPNISLNFYK
						NSVSKLLNRKKGLIHLLQPFILR
						LCVSLHKNREENLGITIODVAM
						RLSALPNIVCRLYKKTASEVLK
						EWSTRWDEYTHHKEVSOKALP
						TLSVKIYKKNICQLLNQNNGPN
						SEMNALITKKSVKSASGYLESI
						AALVEKEISSYKNYTEAFYNLL
						CDVYSHVTEMKLSSIEQIGNTL
						FVESASGVWRALRPRGKEVNDI
		l				FIDQIANSVLVDSSKDICKRFEA
		1				CGAMVKKENLHIKTROKISEKL
						HWPFGAPRLTSLTGPEKTGKGE
	l					HPGAKARGELSGAPPLIHTRKL
						SLMSNVETLLLEKLQGYTGAH
15102	45470	A	15189		966	SEMSIVE TELEBREQUITOAH
15102	45471	A	15190	157	531	GDHTTFLCLLMVNAEFKYNFK
13103	75171	(`	13130	1.57	1551	TIIS*DLLKKDSGOYGHFTIFFRS
						MSIGCFSIYLCIFNFFHQCFAVF
	l					LVEIFTFLVKCIPQVDSIEDTTFG
	i					HPEEATPHPFSFFMRLEOFSHIF
	l	1				SSWPVSFTAH
15104	45472	A	15191	9	494	PTDRPRPRALAGARGPPGRAPR
1.510	1.52			ľ		VPPPAGGPRPSQYGHGGRPNPR
		l				HGWHDAFCPNNTPYR\ERPLDS
	i					RYLTPGPLDMRGDSGFRTGRGS
						DVGSLKPTDTRRNGKIYLNGS/
	l	ļ				NVFYYQQRLHPVHRGDGARRG
	1					FSGRTCLHRMVC*YEQTGHQS
				ŀ		DOT*KARSAYG
15105	45473	A	15192	1361	2038	- (1 19 19 11 19
15106	45474	A	15193	1376	1906	LAFPDGMASLIMMSLPAMRTL
1		1		_		AWAMAVSGNCIPEGQFSALSC
						ADFGISPKLIL*SSSAPN\YLAMG
	l					ISCPRADDLPEAGTAETDRACR
	l	1		l		SIPOYGTDLGWYPATAAGPOSE
	l					SVQGAV*PHRYEAG*ST/VWKP
	l		1			LKKK*KPLPRR*LARWOMIRFR
	l					KPSPLTTSLKLTCA WRFLDFRW
	l					LE
Ц			L			

SEQ ID	SEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon far last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15107	45475	A	15194	h	1388	I MIEHPRRRDNYROSGPPVADKF
15107	13173	l'`	15151	l'		TTKGEPLISDIVSDAOOANLLIP
					1	VDETPPVINDEQSTSTPLTTAQT
						MALAAVADKNTTKDEKADDL
						NEDVTASLSALFAMLPGFDNTP
				1		KVTDAPSTVLPTEKPTLFTKLTS
				1		EQLTTAQPDDAPGTPAQPLTPL
ľ				1		VAEAQSKAEVISTPSPPSISGVT
İ				1		RSKMRQREWLVLPPEDSLVIRR
				i		SNKTSQLPAFNVPLSAAKAAAS
						KPWRASNTACRNSRPSKVCSAE
				l		GEOWFSASGRGEIHSOSPARDM
						AOMCKLAAEGHFAEARVINOR
						LMPLHNKLFVEPNPIPVKWACK
1				i		ELGLVATDTLRLPMTPITDSGR
1						ETVSGDETYLEAAPLAELHAPA
1						GMILPVTSGDYAIPVTNGSGAV
						GSYSAITEPGAGSDVGSLKTTY
						TRRNGKIYLNGS/NVFYYQQRL
						HPVHRGDGARRGFSGOTCLHR
			1			MVC*YEQTGHQSDQT*KARSA
			1			YG
15108	45476	Ċ	15195	1	1905	10
15109	45477	A	15196	285	1706	YARVHSGRSSIRRSIPVSICIRSSI
						RGCVPVVHCIRSSIRRSIPVDHL
						YQIIHTKKHTRCPFVSDHPYEE
1						APDYSASLLPDWLPVPSSKVVI
						TYPGFTSICGYYTTSSKACFSVF
				Į.		AWCFVTSDVPGRFSDEDEDDSE
					1	MWSLTALMGVMGGWKTSARS
						EVSILCSTSSKVIHSSYPYADRA
1				i		FQVWSLDARFAHINKPFGVDRF
						SGEAPSRAITTMYGVKVLNRPA
						TGTAHRLVPPDVPVHPHGGESN
		ŀ				KPRVQTTVSSGISRLSIPISASAL
		ľ	İ			ONVPDKTHDISNITDTVYHCPL
1		1	1	I		SQSSRWYWRFVPVRSPPLSNGQ
		1	1	I	1	TAVRPGGSRAGAAGGPLSEYS
		1	1	I		GSGFGVMKWTKDQLTVTVKV
	1	Ì				HHLPEVVEFLYYKQGGWLSVL
	1		1			FGNDERKLNGHYAVYYVLSRE
	1					KGTKCWI/TDSMDYROR/PAPT
	1	1		l		NDAETYEFINELGARKTTVVPI
ĺ		1		I	1	GPLHVTS/DEPGHLRLFVDGENI
			l	l		IEANNE*IHRTQC
I5110	45478	A	15197	3	1396	
15111	45479	В	15198	1	1399	
		1			L	L

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15112	45480	A	15199	283	796	FVGVKSAGNRLKTIDELANESR
13112	45400	^	13133	203	//	SSVGLTQAALKTLESSGAIRIER
i				1		RGRNGSYLVEMDNKALLTHVD
İ	ŀ					INNVVCAMPLPYTRLYEGLASG
l						LKAQFDGIPFYYAHMRGADIRV
						ECLLNGVYDMAVVSRLAAESY
l	l					LPQKGLCLAL*LGPPPYVGE\YQ
1		i				L*OVKAKSEYVKRVGHV
15113	45481	A	15200	48	3376	- Q.III.III.
15114	45482	A	15201	49	681	DPGADRRGSVWGLSDLRNTYN
						ALYLTHSGRLSECPTPANEEKE
1						PTNHNRTLTNOTOKEKKKTRT
						KDCTTRNKNSTRPGSGIIAGPSY
						YTVAGPPSAIDPRWLRPTPPAL
İ				ł		DPO*MPHTSKRGKRADOSOPN
		l		l		ANKPNTEREKKDKNEGLHHKE
1				l		OEO\ORPGSGIIAGPSYYTVAGP
1				l		PSAIDPRWLRPTPPALDPOTEPL
		l				IFOOLEIDHYVAGLTHEVMSLS
						HLTLVMRWGSVSAAHLAHWA
						RCRWISTELTPNSITSTSSPLKD
						CSSSPAMEQSWLENDFDELTED
						KKTCLCDH
15115	45483	Α	15202	3	587	
15116	45484	Α	15203	3	774	MVLMAGFTAGNEKGELVVLG
		Į.				RNGSDYSAAVLAACLRADCCEI
		l	ĺ			WTDVDGVYTCDPRQVPDARLL
1		l				KSMSYQEAMELSYFGAKVLHP
		l				RTITPIAQFQIPCLIKNTGNPQAP
		l				GTLIGASRDEDELPVKGISNLN
		ı				NMAMFSVSGPGMKGMVGMA
		ı				ARVFAAMSRARISVVLITQSSSE
ŀ		1				YSISFCVPQSDCVRAERAMREE
1		l		1	1	FYLELEESLLEPLAVT\ERLPII\S
		l				VGSDGNPPPCGGISPKFF\AALA
1		l				PRPISKFVALCSGIPF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first eodon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15117	45485	A	15204	1474	1511	HIS-YPANILYMELKACFTGA/NE KGELVVLGHNILAVSYFIALAAS LRADCCEIWTDVDGVYTCDPR QVPDARLLKSMSYQEAMELSY FGAKVLHPRITPIHOGPOIPCIL NTONPOAPCTLIGASRDEDELP VKGISNLNNMAMFSVSGPGMK GMVGMAARVFAAMSRARISV GLITQSSSEYSISFCVPQSDCVR AERAMGEFFVLELKEGLLEPL AVTERLAIISVVGDGMRTLRGIS AKFFAALARANINIVAIAQCSSE RSISVVVNDDDATTGWRVTHQ MLFNTDQVIEVFVIGYGGVGG ALLEQLKRQOSWLKNKHIDLR VCGVANSKALLTNVHGLNLEN WQEELAQAKEPFNLGBLIRLVK EYHLLNPVIDCTSSQAVADQY ADFLREGFHVVTPNKANTISS MDYYHQLRYAAEKSRRFELVE HNTFVSPHIRPLIERRGRSSTRD KWCARPSTERQQDNSARILISC KWCARPSTERQQDNSARILISC KWCARPSTERQQDNSARILISC KQQNITSFEHTISSIRVGTTNSLN
15118	45486	Α	15205	198	462	

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion
15119	45487	A	15206	31	1981	RWLMRGMGIALPAAMASTV
						STIAETPEPPPLSDW/LRASG/D
				l		WVHLLGLDPHLSSLLLSICLF:
						SLIWIF*LOP*SV/ALAFQMGA
				ł		LFASNIGVIEGEGPOAKPL*CL
						VSYFEMLNNLNAQGPKGKLS
				1		COCLSLOVMTMPEYLKKRW
				l		GORSOWPOMHVSFSS*LFPF*
		1				DIFAGAIFIKOPSGEGKYMRT
						G*NALVLFFLSLGGLEVVIINI
	l					CLRFQ*DISFYSDAFLAFNEV
					}	YESFTEKYVNATPSVVEGDN
	j	l		l		ISASCYTPRADSFHIFRDAVTO
						GR*LPLKSSPGLFHECVTFSR
						IVQRCL\CGKDMSHVKAACIN
				1		LFV*WLLVLPIWASLDSLLNS
						DMVACVVPSECVISTLSGLAI
				l		CPATNOCLYPFPAPGLRGLM
						VMLASLMSSLTSIFNSASTRT
		Ì				AQWK*L*CFCLMFKLCCFLLI
						FVLLLTVVSIVWVPLVQVSQ
		!				OLIHYTESI***LT*ISFSYFLNI
		1				FFLLLLQGAFWGLMVGLAM
		l				IRMITEFAYGTGSCLAPSNCPI
		ı		i		CGVHYLYFSIKVPFYYTIANIO
		1				LLMHFCMVPVQLYRLCWVL
		1		1		NTTLTRISSSFVI*IFCNHLLIF
		1				DYPEKSRGCLKKAYDLFCGL
		1				KGPKLTKEEEEALSKKLTDTS
5120	45488	В	15207	1	1392	
15121	45489	Α	15208	57	1523	
15122	45490	Α	15209	473	589	LPRQLALSCVVHCQTLVPAIR
		1			1	DWGWLF*W*SVTPVM

1924

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15123	45491	Α	15210	lı .	1297	MGKIMSSYQHVDLIVDCDFEA
		l				YHROPSSGTKSVVTGPSAQALI
1						TEKIVTYHWDQHVGDVLVLNI
		l				YCVINVLDHTVKAVIDDRFMG
l		l				GITTINLOLMAIIVRYFFASGSA
		1		l		YIVAMLPVFAMLANVSGAPLM
				l		LTALALLFSNSYGGMVTHYGG
		l				AAGPVIFGVGYNDIKSLCFDER
ļ						TRIDVADTYASNWRSRCNQTIF
		l				NGECPNTROHIAAVRRGIDTLL
		1				MNDYLRKQIIDISTRIAGTANN
		1		l		GDFTGQWITAANTVNFQIVAGT
						HNCHQYFIALRLIFGILKYIQDR
		1		İ		AREGLKQRDLAEALSTSPQTVN
		1		i		NWIKRDALSREAAQQISEKFGY
		1		l		SLDWLLNGEGSPKKDLESNIPP
1	į			l		ESEWGTVDAWDKNTPLPDDEV
1				İ		EVPFLKDIEFACGDGRVHDEDH
İ	·	1				NGFKLRFSKV\VQPRVGC*QG*
		1				G*LLYLDSV**GDPLEST
15124	45492	A	15211	62	301	SWMSSPRAFSSSTCLETAAPTR
10.12		ļ.,		1		MATALTSAACRLPPAAGSGSP
		1				WAPAPYRWRARRRTGPMSDR
		l				Q*RTSQPCPEPAAASR
15125	45493	Α	15212	1	1665	MQDSAHLRRLPPAASRRVWLA
l		l				LGSCPLPLESSPAHRAYERPSVA
			l			HQPAVPGARRSLEVPSEVPLTV
ŀ		1		l		PALTPPRIPAMLTAVCGSLGSQ
		1		ļ		HTEAPHASPPRLDLQPLQTYQG
				l		HTSPEAGDYPSPLQPGELQSLPL
		1		l		GPEVDFSQGYELPGASSRVTCE
						DLESDSPLAPGPFSKLLQPDMS
				l		HHYESWFRPTHPGAEDGSWWD
				l		LHPGTSWMDLPHTQGALTSPG
						HPGALQAGLGGYVGDHQLCAP
		1		ŀ		PPHPHAHHLLPAAGGQHLLGPP
				i		DGAKALEVAAPESQGLDSSLD
				İ		GAARPKGSRRSVPRSSGQTVCR
ŀ		1				CPNCLEAERLGAPCGPDGGKK
		1				KHLHNCHIPGCGKAYAKTSHL
		1				KAHLRWHSGDRPFVCNWLFCG
1						KRFTRSDELQRHLQTHTGTKKF
1		1	1	i		PCAVCSRVFMRSDHLAKHMKT
l				l		HEGAKEEAAGAASGEGKAGGA
l			l	l		VEPPGGKGKREAEGSMASSSPD
				l		SPCSCDCFVSVPPASAIPAVIFA
				l		HELGPTPGRGAGGGVCPRRHS
l		1				HSWEPAPGL*AVGGGGGASGQ
		1	١,			AGDRGCHA*EAGRS/SCSRRGS
		1				GL*RAGTS*ISASS
				L	L	1-2

, /=possible nucleotide ssible nucleotide insertion)
HGSSTOTPAHVGSF
D/ACRRSKIRGPGGW
RDGRNSR/DLRNLPC
*AAAPSPPWPAAGA
DGWPHSYSHGHSSA
RSPPLLAGELRPPGIQ
RGILSGSPCPPFSGAG
AMVLL*PPSP/PAWPT
SSKVSSTSRPLRKSIM
RAPTPA*PSASKPTNA
SEPRSHAHLDGRHRD
PLSGPRTLGGKSLRGS
GVPRPRPLPENLDAPL
AGAAVQLA
AHPGGTPREQDGAM
AQAMIWTAGWLLL
QALECYSCVQKADD
KTVKCAPGVDVCT\
TIHGQFSLAVR\GCG
DRGLDLHGLLAFIQL
RCNAKLNLTSRALDP
PPNGVECYSCVGLSR
PPVVSCYNASDHVY
IVTLTAANVTVSLPV
FCTRDGVTGPGFTLS
RCNSDLRNKTYFSPR
PPEPTTVASTTSVTTS
TSTTKPMPAPTSQTPR
SRDEEPRLTGGAAG
GQYPAKGGPQQPHN
AGLAALLLAVAAGV
ECYSCVQKAD/DGC
VKCAPG/VDVCTEA
IPFWIWPRPFNAAPPP
YDTALQARPLY*QL
*ELVH/GQFSLAVRG
K/NDRGLDLHGLLAF
QDRCNAKLNLTS/RG
PPVVSCYNASDHVY/
IVTLTAEPTTVASTTS

SEQ ID NO! Note N	SEO ID	ISEO ID NO:	Met	ISEO ID NO:	Nucleatide	Nucleotide location of last	Amino acid sequence (X=Unknown,
Sequence 99/540/217 codes for peptide sequence deletion, impossible nucleotide insertion) sequence 15130 45498							
VGAQTYNGIELIEVGGNSAGNR ALTIGTVEAGAYVYTLAKGKG NDEKNISPSYTRPEAGSYISMA AANSLISHRLHDRISCLQPLKI LVTGGAGFIGSAVYRHINNTO DSVVAVDKLTYAGNRESLADV SDSERYVFEHADICDAPAMARI FAQHQPDAVMHLAACSHVDRITGPAAFETNIVGTYVLLEAAR NYWSALDSDKNSFRFHHISTE EVYGDLPHPDEVNNTEELPLFT ETTAYAPSSPYSAKASSDHLV RAWKRTYGLPTIVTNCSNNYG PYHPPEKLIPLVILNALE		1			sequence		
VGAQTYNGIELIEVGGNSAGNR ALTIGTVEAGAYVYTLAKGKG NDEKNISPSYTRPEAGSYISMA AANSLISHRLHDRISCLQPLKI LVTGGAGFIGSAVYRHINNTO DSVVAVDKLTYAGNRESLADV SDSERYVFEHADICDAPAMARI FAQHQPDAVMHLAACSHVDRITGPAAFETNIVGTYVLLEAAR NYWSALDSDKNSFRFHHISTE EVYGDLPHPDEVNNTEELPLFT ETTAYAPSSPYSAKASSDHLV RAWKRTYGLPTIVTNCSNNYG PYHPPEKLIPLVILNALE	15120	145.400	 	1.5018		loca.	I DUMODINO CHI TINUNGO
ALTTGTVEAGAVVYTLAKGKG NDEKNSPSYYRPEAGSYISNIA AANSLISHRILIDRISCLQPLKI LVTGGAGFIGSAVVRHINNTQ DSVVNVDKLTYAKGNESLADV SDSERYYFEHADICDAPAMARI FAQHQPDAVMHLAAESHVDRS ITGPAAFIETNIVGTYVLLEAAR NYWSALDSDKKNSFKFHHISIT EVYGDLPHIPEVNTSELLEFT ETTAYAPSSPYSASKASSDHLV RAWKRTYGLPTIVTNCSNNYG PYHFPEKLIPLVILNALE EYYGDLPHIPEVNTSELLEFT ETTAYAPSSPYSASKASSDHLV RAWKRTYGLPTIVTNCSNNYG PYHFPEKLIPLVILNALE AFSON A I5222 I 498 MTRSLLSVFEEDAGTLTDYTNC LQAMQRYYGAQLNLLHTELA KQLLETQWFYLSYNSEKRISQTI SAMWRGRAAQA/TQGHWVWG MVEAVCSKVSTLKDLFGLASN GHDLSMAYSRJFKKKENEKL AAFGQPTWCDPAERVGFLEPV GPDPGAKSAPSPRDRASTL SAMWRGRAAQA/TGGHWVWG GHDLSMAYSRJFKKKENEKL AAFGQPTWCDPAERVGFLEPV GPDPGAKSAPSPRDRASTL SAMWRGRAAQA/TGGHWVWG GROWN GR	15130	45498	A	15217	μ.	964	
NOEKNSPSVYRPEAGSYISNIA				l	1		
AANSLESHRLHDRHSCLOPLX1				i .			
Lytggacpigsavvrhinnto							
SDSERTYTEHADICDAPAMARI				i			
FAQHQPDAVMILARESHVDRS ITGPAAFIETNIVGTYVLLEAAR NNWSALDSDKKNSFRFHISTE EYYGDLPHPDEVNNTEELLFT ETYAYAPSPYSAKASSDHLY RAWKRTYGLPTIVTNCSNNYG PYHFPEKLIPLVILNALE IS132 45500 A 15222 1 498 MTRSLLSVFEEDAGTLTDYTNC LLQAMQRVYGQAUNLHTELA KQLIETQWFYLSYNSEKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT SAMWRGRAGA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT SAMWRGRAQA/ATGGHWWG MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKRISOTT 15134 45500 A 15223 1 1244 15154 15134 15235 1 1404 1413 145509 A 15233 1 1404 1413 1414 145509 A 15233 1 1404 1415			ł				
ITGPAAFIETNIVOTYVLLEAAR			1		Ì		
N.W.SALDSDKKNSFRFHHISTE							
EVYGOLPHPDEVNNTEELPLFT ETTAYAPSSPYSASKASDILV RAWKRTYGUPTVTNCSNNYG PYHPPEKLIPLVILNALE			1				
Section							
			1		ľ		
PYHFPEKLIPLVILNALE	i						
	1						
15132 45500							PYHFPEKLIPLVILNALE
LLQAMQRYYGAQLNLLHTELA KQLLETQWFYLSYNSEKRISQTI							
	15132	45500	Α	15222	11	498	
SAMWRĞRAAQA/TQGHWVWC MVEAVCSKVSTI.KDLFGLASN EHDLSMAKYSRI.PKKENERL		1	1		İ		
MVEAVCSKVSTLKDLFGLASN EHDLSMAKYSRLPKKKENEKL AAFQQPTWCDPAERVGFLEPV GPDPGAKSAPSPRDRASTL IS133 45501 A 15223 1 1224 IS134 45502 A 15224 1 1353 IS135 45503 A 15225 1 2580 IS136 45504 A 15225 1 1260 IS137 45505 A 15226 1 3117 IS138 45506 A 15228 1 2142 IS139 45507 A 15229 1 1413 IS140 45508 A 15230 1 4049 IS141 45509 A 15231 1 381 MLMLNIGFKNFQFEHLKCSHGS YSEDIGFPNAGALSGAMFLLEE KHLLPGLKEEERLERASIRESQ QRRMERKOHQRGLSASYUWL DTMRRKARSPYSLAAIKNFY KGGIRDQKPSYDSLNLCMT KGGIRDQKPSYDSLNCMT KGGIRDQKP			1		1		
BIDLSMAKYSRLPKKKENEKL AAFGQPTWCDPAERVGFLEPV GPDPGAKSAPSPRDRASTL 15133 45501		i .					
Section			1				
Section	1		1		ŀ		
15133 45501		i			ł		
15134 45502	15100	45501	ļ.,	15222	ļ. — —	1224	GPDPGAKSAPSPRDRASTL
15135 45503 A 15225 1 2580							
15136 45504 A 15226 1 3117							
15137 45505							
15138 45506 A 15228 1 2142							
15139 45507 A 15229 1 1413							
15140 45508							
15141 45509 A 15231 1 381 M.M.L.NIGFKNFOFEHLKCSHOS YSEDIGFPNAGALSGAMFLLEE KHLLFGLKEERLRASIRRESQ QRKMKEKQHQRGLSASYUWAL DTMRRKARSPYSLAAIKNEY KGGIRDQKPSYDSLNLCMT 15142 45510 A 15232 1 467 DHSSFAREQNWMENEFDELT EVGRRWVITNSELKKHVLTQ CKEAKNLERRLGELLTRITSLE KNINDLMELKNTARELHEAYTE INSWINQAEERLSEIGDQLNEIK REDIKITEKMNSTITVRVAASM QSKLLQGVAEEGPLRLTRASSF 15143 45511 A 15233 1 1140 1141							-
YSEDIGFPNAGALSGAMFLLEE KHLLFGLKEERLRASIRRESQ QRKMEKQHQRGLSASYWML DTMRRKARSPYSLAAIKNEY KGGIRDQKPSYDSLNLCMT IS142 45510 A 15232 1 467 DHISSSPAREQNWMENEFDELT EVGFRRWVITNSSELKKHVLTQ CKEAKNLEKRLGELLTRITSLE KNINDLMELKNTARELHEAYTS INSWINQAEERLSEIEDQLNEIK REDKITEKMKNSTYTVRVAASM QSKLLQGVAEEGPLRLTRSASF 15143 45511 A 15233 1 1140 1144 45512 A 15233 1 2640 15145 45514 A 15235 1 714 15146 45514 A 15235 1 714 15146 45515 B 15237 1 735 1 735							MLMLNIGEKNFOFEHLKCSHGS
KHILIFGIKEERILRASIRRESQ QRRIMEKQHQRGLSASYUWIL	10.1.1	1.000	Γ.		ľ		
	1		ł				I .
DTMRRKARSPYSLAAIKNRY KGGIRDQKPSYDSLNLCMT		1	ŀ				
KGGIRDOKPSYDSLNLCMT	1						
15142 45510			1				
EVGFRRWVITNSSELKKHVLTQ CKEAKNLERILGELLTRITSLE KNINDLMELKRTARELHEAYTE INSWINQAEEKLSEEDQLAEIK REDKITEKMKNSTTVRVAASM QSKLLQOVAEEGPLRLTRSASF 15144 45512 A 15233 1 1140	15142	45510	A	15232	1	467	
CKEAKNLEKRLGELLTRITSLE	1.5172	1.55.0	()		ľ	I	
KAINDLMELKNTARELHEAYTS INSWINGAERLSEIEDQLNEIK REDKITEKMKNJAGERLSEIEDQLNEIK REDKITEKMKNSTYTVRVAASK QSKLLQGVAEEGPLRLTRSASF 15144 45512 A 15233 1 1140 15144 45512 A 15235 1 714 15146 45514 B 15236 1 546 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 15147 45515 B 15237 1 735 1 735 1 73516			1				,
NSWINQAEERLSEIEDQLNEIK REDKITEKMKNSTTVRVAASM QSKLLQGVAEEGPLRLTRSASF			1				
REDKITEKMKNSTTVRVAASM QSKLLQGVAEEGPLRLTRSASF 15143 45511							
QSKLLQGVAEEGPLRLTRSASF QSKLLQGVAEEGPLRLTRSASF	1	I	l	1	l	1	
15143 45511 A 15233 1 1140 15144 45512 A 15234 1 2640 15145 45513 A 15235 1 714 15146 45514 B 15236 1 546 15147 45515 B 15237 1 735				l	1	1	
15144 45512 A 15234 1 2640 15145 45513 A 15235 1 714 15146 45514 B 15236 1 546 15147 45515 B 15237 1 735	15143	45511	A	15233	1	1140	(3(2 <u>2.0.</u> 2
15146 45514 B 15236 1 546 15147 45515 B 15237 1 735		45512	A	15234	1	2640	
15146 45514 B 15236 1 546 15147 45515 B 15237 1 735			A	15235	1	714	
15147 45515 B 15237 I 735			В	15236	1	546	
15148 45516 B 15238 50 1603	15147	45515	В	15237	1	735	
	15148	45516	В	15238	50	1603	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15149	45517	A	15239	1374	2220	QAENSHININKDVHISETPSEG HORKRRKDHSSSPAEGNWE NEFDELTEVGFRRWVITNSSEL KKHVLTQCKEAKNLEKRLGEL LTRITISLERNINDLMELKNTAR ELHEAYTSINSWINOAEERLSEI EDQLNEIKREDKITEKMKNSTT VLVSTWGSQLEQNNLPHISPLF HSTHDLNISPYGGCLLQACSSK LQGLAEGGPRLTRSASFFAA KLCLTKHFLPTEMILIKSRKTP DTOPRRLYLLQGVVQTISRKG PTALNIRGWKTFAAAPPIRK
15150	45518	В	15240	1	1311	
15151	45519	В	15241	46	212	
15152	45520	A	15242	3	371	EPLPPEPPAVPSEVELQHLRKEL ERVAGELQSQVKNNQHISLLNR RQEERIREQEERLRKQEERLQE QHEKLRQLAKPQSVELK\SQEV QSLQQQPDHF\LGHL\QQYV\AT LSAARWAAYQQLD
15153	45521	Α	15243	392	558	
15154	45522	В	15244	52	689	
15155	45523	В	15245	400	603	
15156	45524	А	15246	5	1158	RKFWRWMVVMAAQHSQAAQ QQRR**VSKGP**PGPMGGPVL GHWWHSGGISLAVPSLPPPGKS SVFLFLFSASFSDMKLNPGTVSP RLIFGSYDDGFLVETVVKFDFQ LWQKTPLHRAAGERRRAEQRG KLSFVMP

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15157	45525	A	15247	1	1808	MAEETQHNKLAAAKKKSATGF
		1		1		HREGPTSSATLKDLESPCOERA
				l		VVLDSTSVKISRLKNTIKSLKQQ
		1				KKQVEHQLEEVTAPDNLVPWV
				l		GLSWGIGGILGACLLLCHLCLP
				l		LEKKANNEROKAERELEVOIOT
				ĺ		LIIOKEELNTDLYHMERSLRYFE
				.0.		EESKOLAVRLOHSLOCKGELES
		1	l			ALSAVIATEKKKANOLSSRSKA
	1					RTEWKLEQSMREEALLKVQLT
		ļ.				QLKESFQQVQLERDEYSEHLKG
		ľ		1		ERARWOORMRKMSQEICTLKK
	1					EKQQDMRRVEELERSLSKLKN
						OMAEPLPPEPPAVPSEVELOHL
						RKELERVAGELQSQVKNNQHIS
	ĺ					LLNRROEERIREGEERLRKGEE
		1		1		RLQEQHEKLRQLAKPHCAFSRS
						*TMRTRAHLQLEQQVKELQE/Q
						ALARVFLQVELKSQEAQSLQQ
				i		QPDHYLGHPAAEHLEAASQQN
	l	1				QQLTAQLSLMALPGEGHGGEH
	1					LDSEGEEAPRPMPSVPEDLESR
1	1					EAMVAFFKSAGASAQEKQAQL
1	1					QEQVKEQRVCCQRLAHPVASA
						QKEPEAARGPGAPGPGGESSSF
	į.			l		MDHLKEKADLSE/PGEKRTLLH
						P/PTG/GDRRHQKTHHLLSEPGG
	1	1				CAK/DAALGPGHHQAGAQGGD
15158	45526	Α	15248	2085	2700	YPGKRGLEWT*ANSNRPAAEG
						PDE\SSSLPATEQSWMENDFDE
i		ĺ				LRKEGFRRSVITNFSKLKEDVQ
1						THRKEAKNLEKGLDEWLTRINS
1	1	1				IEKTLNDLMEMKTMA*ELHDT
						CTSFSSRVDQVEERLSVI*DQM
ı	1	1				NEMK*EEKFREK\RVKRNEQSL
1						QEIWNYVKRPNLHLIGVSEIDR
1						ENGTKLENTLQDIFQENFTYLA
		1				RQANIQIQEI

NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15159	45527	A	15249	378		QTERNSININKADIHTKTPSVRH HHQRPKLIKITKIMGKKIJSKA ENSKNQSGSLPPKECSSSPAVE QSWTENEPDEFTEGEFRRIESQ QEIWDYVKRPDLRLIGVPDSDG ENGTLENTLQJITENEPPILAR QASIQJEIQWTOKYSSRRATP RHIVRFTKVEMKEKMLRARE KELKIKKLTQNCLTTWKLINIL LNDYWYHNEMKGEIKMFFETH ENIDTTYQNLWHTFKAVCRIK FIALNAHKRKQERSKIDTLASQ LKEIEKQEQTHSKASRRQEITKI RVELKEIETQSSLQKINESRSWF FEKINKIDRPLPRLIKKREKNQ YYKHLYANKLENLEEMDKFLD TYTLSGLNKEEVESLNTISTIGSE VYAHLYANKLENLEEMDKFLD TYTLSGLNKEEVESLNTISTIGSE VALINSIPTIKKSRTRWHSRIL
15160	45528	А	15250	935	1269	PEVQGGAA LIEGKLTERNSININKKDIHTKT PSVG/PPSSKTKDRSSLPAIEQS WMENDFDELTEVGFRRSVITNL SKLKEDVRIPHKEATNLEKRLD EWLTRINSIEKTLDDLMELKTM
15161	45529	A	15251	401	497	KTFSPFYFS*RRRTVRDINGFTE Y*EPEKLLG
15162	45530	В	15252	220	940	
15163	45531	Α	15253	1	1476	

1930

SEO ID	SEQ ID NO:	Mat	SEO ID NO:	Nucleotide	Nucleatide Idention of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
10100		<u> </u>	10004		2423	NO CONTROL OF THE CONTROL OF THE
15164	45532	Α	15254	1	2423	MNGKNKTNIYQEPTVCQALHW
		1				TLARIGKEEPRRGQRGEGLLCS
		l		i		RERQPRGLSSPGTAAPAAPAAP
		l	l	1		TPPALPGHRTRSGSPEGPEPILR
		l				RGGSADAPGRVKTWVNRPAAF
						VGKCLGDGSADTACDQAMKG
ł		l				QRGSNRKRKTSHSLTFPTSSDM
		1				HAGGLSAAHPCAHSRAPPPASP
		ı				SPYKGSLGTRTDGLTDSRADSA
		1				ASGARALGVDGQPQPSPPRRAF
		1	l	1		PRPVGADGSSRGPQPGRRRVER
						RGPATQPGDSRALPEPRGVPAV
		l				HPAGSGSEWERPPPAAPSPEHR
						DKMLPGLRRLLQASARRPQPFP
						EAARWVWSPCFSSGTLCPSFQS
		1				ERPLLPFQGASFPGTTLHQRLCF
						VAALHGSGKGQKWQQSHCYA
					1	AAVMEEPPAPGRSSESLOGPLS
						WAGGAPASACLLLMLLALPLA
						APSCPMLCTCYSSPPTVSCQAN
						NFSSVPLSLPPSTQRLFLQNNLI
						RTLGARHLWVQPAHPVALLQQ
						PLHHLPGHFPPL\QALEELDLGD
		1				NRHLRSLEPDTFOGLERLOSLH
		l			\	LYRCOLSSLPGNIFRGLVSLQYL
						YLOENSLLHLQDDLFADLANLE
	l		Į.			PTSFLHGNRLRLLTEHVFRGLG
						SLDRLLLHGNRLOGVHRAAFR
		1			\	GLSRLTILYLFNNSLASLPGEAL
		1				
						ADLPSLEFLRLNANPWACDCR
		1				ARPLWAWFQRARVSSSDVTCA
		_		10	100	TPPERQGRDLRALREADFQACP
15165	45533	B	15255	47	482 636	MRDPNTKRSRGFGFVTCATVE
12166	45534	l ^A	15256	ľ	030	
1		1	1	1		EVDAAMNARPCKVVGRTVEPK
		1	1			RAVSREDSQRPEDTEELHLRDY
		1				FEQYGKIEVIEIMTDQGSGKKR
l	1	1	I	1		GFAFVTFDNHDSMDKTVIQKY
		1	1			HTVNGHNCEARKALSKQEMAR
		1	1			ASSSQRGRSGSGNFGGGRGGGF
		l	1			GGNDNFGRGGNFSGHGGFGGS
1		1	1			HGGGGYGG/SGDGYNGFGNDG
	1	1	1		1	GGGSYNDFVNYNNQSSHFGPM

	O 01/0/5067 PC 1/US01/08631							
SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,		
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)		
				sequence				
15167	45535	A	15257	1	890	MSKSESPKEPEQLRKLFIGGLSF		
15101	10000			ľ		ETTDESLRSHFEQWGTLTDCVV		
	İ					MRDPNTKRSRGFGFVTYATVE		
						EVDAAMNARPHKVDGRVVEP		
						KRAVSREDSQRPDYFEQYGKIE		
						VIEIMTDRGSGKKRGFAFVTFD		
						DHDSVDKTVIQKYHTVNGHNC		
	ŀ					EVRKALSKQEMASASSSQRGRS		
				1		GSGNFGGGRGGGFGGNDNFGR		
						GGNFSGRGGFGGSHGGGGYGG		
İ				1		SGDGYNGFGNDGSNFGGGGSY		
1						NDFGNYNNOSSNFGPMKGGNF		
1						GGRSSGPYGGGGQYFAKPRNQ/		
						GGYGGSSSSSSYGSGRRF		
15168	45536	A	15258	1	286	00100333333103000		
15169	45537	A	15259	1	1674			
15170	45538	A	15260	235	1370	KREGEGRLADTFVSLLLSARGR		
13170	75550	^	13200	1222	1370	RRRSIVKVSLHPAVMSKSESPK		
						EP\EQLRKLFIGGLSFETTDESLR		
						SHFEQWGTLTD\CVVMRDPNT		
1				l		KRSRG\FGFVTYATVEEVDAAM		
1						NARPHKVD\GRVV\EPKRAVSR		
1						EDSORPGAH\LTVKK\IFVGWH*		
1						RKTTFWGGKKKRAKKHHLRD		
1				1	İ	YFEQYGKI\EVIEIH*LDR\GQWP		
		i		ì		RKRGFAFVTF\DDP\DSVDKIVI		
			ĺ	i		OKYHTVNGHNCEV\RKALSKO		
						E\MASASSS\QRG\RSGSGNFGG		
						GRGGGF\GGN\DNFG\RGGNFS\		
						G\RGGFGGT\RGGGGYGG/SVG		
						DGYNGFGNDGSNFGGGG\SYN		
						DFGNYNNQ\SSNFGPMKG\GNF		
						G\GRSSGPYG\GGGGYF\AKPR\N		
						OGGYGGSSSSSSYGSGRRF		
15171	45539	В	15261	1	660	Q00100333331030R0		
15172	45540	A	15262	2	486	KMRQRERAQGKK/VAPAPAVV		
1.51,72	1.5540	ľ		Γ	1	K/KQEAKKVVNPLFEKRPKNFG		
						IGODIOPKRDLTRFVKWPRYIR		
						LORORAILYKRLKVPPAINOFT		
						OALDROTATOLLKLAHKYRP/G		
						DKGALAKLVEAIRTNYNDRYD		
	l			1		EIRRHWGGNVLGPKSVARIAKL		
				1		EKAKAKELATKLG		
1				L		CRARARELATREU		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15173	45541	Α	15263	13	845	PPKMPKGKKAKGKKVAPAPAV
						VKKOEAKKVVNPLFEKRPKNF
						GIGQDIQPKRDLTRFVK\WPRYI
					İ	RLOR\ORAILYKRLKVPPAINOF
						TOALDROTATOLLK/LVAHKYR
						P\ETKQEKKORL\LARA\EKKAA
		l				WQKGTSPTKRPP\VLR\AGS*HP
i		l				VTTLVE\NKKASAGWWIAHRR
1		l				GFPSSLVVFLP\ALCREKWGSPY
1		l				\CIIKGKARLG\RLVTRKTCTTV
		1				AF\TQVNLGRQRALLLKLVEAI
		1				RTNYNDRYDEIRRHWGGNVLG
		1				PKSVARIAKLEKAKAK\ELATK
15174	45542	A	15264	1	5796	
15175	45543	A	15265	1	2212	MPRVEKNIPGRRNSKGFEDPLE
						VTQDVTREWAKKVVWKREKA
1						SKINGAYFCEGRVRGEAIRIRT
1			İ			MKMRQQATLTMTVDKGDNVN
		1				ISFKKVLIKEEDAVIYKNGSFIH
1	1	l				SVPRHEVPDILEVHLPHAQPQD
						AGVYSARYIGGNLFTSAFTRLI
						VRILPPQPENIKISNITHSSAVIS
		l				WTILDGYSISSITIRYKVQGKNE
		l				DQHVDVKIKNATITQYQLKGLE
						PETAYQVDIFAENNIGSSNPAFS
		1				HELVTLPESQAPADLGGGKML
		1				LIAILGSAGMTCLTVLLAFLIILQ
		1				LKRANVQRRMAQAFQNVREEP
		1				AVQFNSGTLALNRKVKNNPDP
1						TIYPVLDWNDIKFQDVIGEGNF
		1				GQVLKARIKKDGLRMDAAIKR
						MKEYASKDDHRDFAGELEVLC
		1				KLGHHPNIINLLGACEHRERGD
		1				RRHLENQVTPTLILRFSNTLSKR
		1				HTRRLYPEPGSEGPTPTEPRSLL
		1			1	AQQSEIKLHGGSEPPLLIFRQTG
		1				SGVDLQQTPTNLQLRVLTVRR
		1	İ			KTDKRKGHPHQKPICSSPSSKT
1		1				KEGFRRSNYSKLKEEVQTHGK
				I		EVKNLEKKLDKWLTRITNAEK
		1				SLKDLMELKTKARELCDECASL
1		1		l		TSQFDQLDKRVSVMEDQMNE
				I		MKQEEKFREKRIKRNEQSLQEI
						WDYVKRPNLCLIGAPESDGEN
1		1				GTKLENTLQDIIQENFPNITRQA
				I		NIQIQEIQRTPQRYSSRRATPRHI
		_				

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15176	45544	A	15266	1	743	PPLLIFROTGSGVDLOOTPTNLO
15170	1.55.11	ľ.		ľ		LRVLTVRRKTDKRKGHPHOKP
		1				CSSPSSKTKEGFRRSNYSKLKEE
	1					VOTHGKEVKNLEKKLDKWLTE
						ITNAEKSLKDLMELKTKARELC
						DECASLTSQFDQLDKRVSVME
						DOMNEMKQEEKFREKRIKRNE
i		1				OSLQEIWDYVKRPNLCLIGAPE
		1				SDGENGTKLENTLODIIOENFPN
						ITRQANIQIQEIQRTPQRYSSRR
	1					ATPRHIIVRFTKVEMKEK\MLR
						AAREK
15177	45545	A	15267	1	618	AAREK
15178	45546	A	15268	200	618	EVSRNNTRTKTSYKODMNTSK
13170	133.10	ľ.	13200		0.0	GF*KDTST*PVT/KERSSSPATEO
				1		CWTENDFDELREEGFRRSNYSE
		i				LKEEVRTNGKEVKNLEKKLDE
Ì	1					WLTRITNAEKSLKDLMELKTM
ŀ						ARELCDERTSLSSRCDQLEERV
		1				SVMEDEMNEM
15179	45547	A	15269	158	594	SGSSICCSPISTVLOPPLLTPROT
13175	143547	l^	1320)	1.50	334	GSGATSSK/SPTDLQLRECSSSP
						AKEQSWMENDFDELREGFRRS
	1	1				NYSELKEEVOTHGKEVENLEK
	1					KIDEWLTRITNAEKSLKDLMEL
	1	1		ŀ		KTMAREOCDECTSLSSRRDOLE
1		1		ı		ERVSVIEDQINEMK
15180	45548	Α	15270	3	1453	LSDLELAAQSIIFIFAGYETTSSV
1.0.00	1.00.10			ľ		LSFTLYELATHPDVOOKLOKEI
	i				· ·	DAVLPNKERSSSPATEQSWTEN
	i	İ				DFDELREEGFRRSNYPELKEEV
ļ						RTHGKEVRNLEKKIROM\VTRII
						NAEKSLKDLMELKTMARELRD
						ECTSLSSQFNQLEERVSVMEDQ
						MNEMKKEEKFRRKRIKRKEOS
						LOEIWDYVKRPNLRLIGVPESD
						GENRTKLENTLQDIIQENFPNLA
						RQANIQIQEIQRTPQRYSSRRAT
						PRHIIVRFTKVEMKEKMLRAAR
						EKEIOTTIREYYKHLYANKLEN
	i					LEEMDKFLDTYTLPRLNOEEVE
	1					TLNRPITGSEIEAIINSLPTKKSP
l		1			1	GPDGFTAEFYQRYKEDLVPFFL
				1		KLFQSIEKEGILPNSFYEASIILIP
l		1			1	KPGRDTTKKENFRPISLMNSDT
l		1			1	KILNKILANRIQQHIKKLIHHDQ
	1	1				VGFIPGMQGWFNIRKSINIIQHI
l		1				NRTNDKTHMIISIDAEKAFNKIO
						QPFMLKTLNKL
15181	45549	c	15271	1111	269	VI I INDIK I DINKE
13191	43349	1	13271	pro	1207	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
15182	45550	Α	15272	94	4505	YLDMMSFVQKGSWLLLALLHP
		1		l		TIILAQQEAVEGGCSHLGQSYA
		i				DRDVWKPEPCOICVCDSGSVLC
		l		l		DDIICDDOELDCPNPEIPFGECC
		l				AVCPOPPTAPTRPPNGOGPOGP
		1				KGDPGPPGIPGRNGDPGIPGQP
		1				GSPGSPGPPGICESCPTGPONYS
		1				POYDSYDVKSGVAVGGLAGYP
		1				GPAGPPGPPGPPGTSGHPGSPGS
						PGYQGPPGEPGQAGPSGPPGPP
		l				GAIGPSGPAGKDGESGRPGRPG
						ERGLPGPPGIKG
15183	45551	A	15273	1628	1832	RRRRKARERRRLAAEQGRRMK
		l				EOCKDGFSE/OASGALKLMGSN
		l				EGEFKAEGNSKFTYTVLEDGCT
		l				KHTG
15184	45552	Α	15274	1	362	
15185	45553	Α	15275	1076	1240	LRQLL*LCPSWRLCLLGGWQL
						HLFF*RHHFVHLVSQRCSIAMS
						PSPRRCFSLHL
15186	45554	В	15276	480	1167	
15187	45555	Α	15277	3	379	
15188	45556	В	15278	13	369	
15189	45557	Α	15279	112	1112	WPPASPSASVIRTVKEFALTNPE
		l				KSSTKETERKETKAEEELDAEV
		l				LEVFHPTHEWQALQPGQACPC
		1		1	1	RNPTYGLNLQTGEREAKLQYE
				1		DKFRNNLKGKRLDINTNTYTSQ
	1	1				DLKSALAKFKEGAEMESSKED
		ı				KARQAEVKRLFRPIEELKKDFD
		l				ELNVVIETDMQIMVRLINKFNS
	1	l				SSSSLEEKIAALFDLEYYVHQM
		l				DNAQDLLSFGGLQVVINGLNST
		İ				EPLVKEYAAFVLGAAFSSNPKV
		1		1	1	Q/EWEAIEGGALQKLLVILATE
				1	1	QPLTAKKKVLFALCSLLRHFPY
1		1		1	1	AQRQFLKLGGLQVLRTLGAGE
	1	1			1	GHGGARRARGHTALRPCHREK
	1					CSPRRRLS
15190	45558	В	15280	54	385	
15191	45559	В	15281	274	776	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
15192	45560	A	15282	362	1323	HQLGRSALLDTRTVSMQKAKN
						GDAVTEPOVA/ENEANGKDTA
1						EVDLLTKELEDFEMKKATARA
		l				VTGILASHPCSTDVHIINLSLTF
						HGQELPSDTKLELNSVRHYGLI
						GLNGTENSMLLSAVGKHGVPIP
						EHIDIYHLTREMPPSDKTPLHSL
						VTVSSPRQRDDMSDGHLCTAR
						RHTDIRDDRSPDAPDTARSLAR
l						STITTRRIFALATPSRTLQQVCS
		ŀ				ALAYTSHPAARTOPLASTALTR
		l				RYMLPSPQLPSLRDHRRTNPLH
						SAVATIPHTTTGEADVSLLCIGL
i		1				STLHYALVYHVSVLTALSAQS
1						MSATDASIIFFEPKR
15193	45561	Α	15283	3	537	
15194	45562	Α	15284	1	455	MPRPEKLGKTQVPPLAECLTKD
1		l				SFLGQKQASLGPLTFRDVTIEFS
	1					LEEWQCLDTVQQNLYRDVMLE
		l				NYRNLVFLVISS/HFTQDFWPD
		1				QSIKDSF/QRIILRTYARCGHKN
						LRLEKIVKVSMRKHQIIHTGKK
		1				PYKCEECAKLLASPQPLENMR
15195	45563	Α	15285	1	397	FAVLMAHYDVQEEDPVLTVIT
		1				YMGLSVSLLCLL\LAALTFLLC
1		1				KAIQNTSTSLHLQLSLCLFLAHL
	1					LFLVAIDQTGHKVLCSIIAGTLH
		l				YLYLATLTWMLLEALYLFLTA
		_				RNLTVVNYSSINRFMKKLMFPV
15196	45564	Α	15286	191	1332	AIETQAITDNCSEERKTFNLNV
		1				QMNSMDIRCSDIIQGDTQGPSVI
						AFISYSSLGNINATFFEEMDKK
						DQVYLNSQVVSAAIGPKRNVSL
		ľ			1	SKSVTLTFQHVKMTPSTKKVFC
		ŀ				VYWKSTGAG/CSQWSRDGCFLI
				1		HVNKSHTMCNCSHLSSFAVLM
				1		ALTSQEEDPVLTVITYVGLSVSL
						LCLLLAALTFLLCKAIRNTSTSL
						HLQLSLCLFLAHLLFLVGIDRTE
1						PKVLCSIIAGALHYLYLAAFTW
						MLLEGVHLFLTARNLTVVNYS
						SINRLMKWIMFPVGYGVPAVT
1				I		VAISAASWPHLYGTADRMLAF
1		1		I		KATAQLFILGCTWCLGLLQVGP
1	1	1		I		AAQVMAYLFTIINSLQGFFIFLV
1			l	1		YCLLSQQAKHKFTPMEVPRQN
		<u> </u>		2.0		HRCLQQN
15197	45565	A	15287	263	397	GRPPAPAARVSGPASVSAG*KR
	l	1				VSASGSVGVRSMRRGSALLPPN
	L	1				F

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino neid sequence (X=Unknown, "=Stop codon, !=possible nucleotide deletion, !=possible nucleotide insertion)
15198	45566	A	15288	1	278	MKFHDRGQMQPKPSPAEAVNS APQREPEGSILRAADPLVWAAK AMSASR/RGS/CGGSSADPRHTE RTPTEPDADTRFQPAETLAGPK TRAAGAG
15199	45567	A	15289	5	545	SEMTEAKSSPNPAQORLSTORP SESRKVPSSELQTLSCGLQRPCL HPGRPGPDPYYASASTRTRSPS ARHAVWGRRGCYPEA/AVRAR EGPHPTLAPASTYGLASPPRAD SAHRPRHRSLTRLLPNRKLGGS SADPRHTERTPTEPDADTRFQP AETLAGPKTRAAGAGGQPWDS ANASPP
15200	45568	В	15290	121	822	
15201	45569	A	15291	1	1308	MI ENVGIEISNGOWSTPPCQVL HLEGNDIYNPKEQGLGNTCRW GWMAAGTRENLCWAPPPENP VASLVLHSTIQIALQESKORVL GTLINSPLVTSTDKANWSLRLA LEVGDOGGLVGLSPPVGSGA MSRCPNCTRKLIGSGISPTSSTL AEEQGSVVKTHSHQGGARCSK EKGWQILLILVAGLCQLVVVV ONVGYKRIPRKYLAALRVAIGH ALAIERRRIRVDVADQAQDTDR DDRSFGRDRNRDSDKTDTDWR ARSATDSFDDYPPRRGDDRFGD KYRDRYDSDQVGRGYWYEYR DGPCRDMDRYGIGGSGRRAFG SGY/RRNDVRGGRDRYEDRY NRWDDGSCSSRDDFSRDDCRY NRWDDGSCSSRDDFSRDDCRY NRWDDGSCSSRDFSRDCRY NRWDDGSCSSRDFSRDCRY NRWDGSCSSRDFSRDCRY NRWDGSCSSRDFSRDCRY NRWDGSCSSRDFSRDCRY DRAPREPRENLILKPHNTPKGD DSSASTQSSRASFVFGGAKPV DRAPREFRYERLILKPHNTPKGD DSASTGSSRASFVFGGAKPV DRAPREFRYERLILKPHNTPKGD DSASTGSSRASFVFGGAKPV DRARAETKVEEWLQKEQDKLQ HQLDEAKLE
15202	45570	Α	15292	2	293	
15203	45571	С	15293	56	298	
15204	45572	A	15294	1	273	TRGPWCDSVLRGCSLEQRSFIS VRLLSYLSACRHPMEDSMDMD MSPLRPQNYLFG\SLGAGAKDE LHIVEAEAMNYEGSPIKVTLAT LKMS

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleoride
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15205	45573	A	15295	3	1024	CGGIHSVLRGCSLEPAFFYLRSA
13203	43373	^	13293	ľ	1024	FSPTISACRHPMEDSMDMDMSP
						LRPQNYLFGCE\LKADKDYHFK
						VDNDENEHQLSLRTVSLGAGA
						KDELHIVEAEAMNY\EGSPIKVT
						L\ATLKMS\VQPTVFPWGAFEIT
1						PTQWSLRLKCG\SGPVHI\SGQH
						FSKLLEEDAE\SEDEEE\EDVKL\
						LSISGK\RSAP\GGGSKVPOKKS
						KTLLLMKDDDDDDEED\DDED
						DG*WMDFGWMEEAEEKAPSE
					1	RNPIRDTPAKNAQKSNQNGKD
		l				SKPSSTPR\SKGQESFKKQEKTP
						KTPKGPSSVEDIKAKMQASIEK
						GGSLPKVEAKFINYVKNCFRMT
						DQEAIQDLWQWRKSL
15206	45574	В	15296	1	1107	
15207	45575	Α	15297	1	1968	MLASMGVVAKEMKYLSSPYPQ
						HAFHTRIKSLMTVNFQAQLWL
						PNEAGPTGDGDWGGDTHGCPS
					1	FSPHLPRFIQKQPWGTGRRLLQ
						EHFSCKSLVPVIPSRDIEAFHLH
						ADLILNTLPSLVIIIGQIESHSEEL
						RARVSTYESGEDTIQCITVTWK
						PKYGATRWKDPGSPDGPLEER
1						RLEDLSTRNNSIRLFRGCGYRV
			ĺ			KQNSSQRPNKVCGLPDTGVLL
		l				QTKCGLVRNSQRSAQTNTKQH
1						EELFSTAGIPENLSRRQEVLPGH
					1	GFTGWVGWTWKLTSGEGVTF
						HHLSATHKLSHASCRQGNFKP
						KGAMSVTRSTVHPHLAFLSAEI
1						NGKCGRPCVFLGIWAINCGMAI
		ļ			İ	LYERLSALVCVSVSERHLDYT
1						MRFLFDIPPSTLEARIGSPLWAL
						SLHPLRSSTVTSSVELILPECTV
1						AVLEIQYVFLGDTILFTTVGKA
				1		RSGTRDRDGGRAQIMQVLQGK
				l		GRELDAAAQPEGQLLREVRVL
		l	1	l	l	GVPFIPRARVDAWLVHTVAVG
				1		SADEAHGLLGAAAASSTGGAG
				l		ASVDGGSQAVQGGGGDPRAAR
				1		SGPLDAGEEEKAPAEPTAQVAD
	i			l		AGGCASEENEVLREKHEAVDH
				l		SSQREENEERVSALKENSLQQN
				1		NDDENKIAEKPDWEAEKTSESR
L			L	L	L	NERHLNGADTSFF\SLEDLFQLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15208	45576	A	15298	1	2104	MKHLKRWWSAGGGLLHLTLL
				Ī		LSLAGLRVDLDLYLLLPPPTLL
						QDELLFLGGPASSAYALSPFSAS
1						GGWGRAGHLHPKGRELDPAAP
						PEGOLLREVRALGVPFVPRTSV
						DAWLVHSVAAGSADEAHGLL
	ŀ					GAAAASSTGGAGASVDGGSQA
						VOGGGGDPRAARSGPLDAGEE
						EKAPAEPTAQVPDAGG\CASEE
	İ					NGVLREKHEA\VDHSSOHEENE
						ERVSAQKENSLQQNDDDENKI
						AEKPDWEAEKTTESGNE\RHL\
						NGTDTSFSL\EDLFOLLSSOP\EN
						SL\EGISIWGDIPLP\GSISDGHEF
						FKHIYHVNFQPRLIS\QD\VNLH
i	İ					EAILLCPNNTFRRDPTARTSOSO
						EPFLQLNSHTTNPEQTLPGTNLT
1						GFLSPVDNHMRNLTSODLLYD
1						LDINIFDEINLMSLATEDNFDPI
	i					DVSQLFDEPDSDSGLSLDSSHN
						NTSVIKSNSSHSVCDEGAIGYCT
				1		DHESSSHHDLEGAVGGYYPEPS
						KLCHLDQSDSDFHGDLTFQHVF
						HNHTYHLQPTAPESTSEPFPWP
						GKSOKIRSRYLEDTDRNLSRDE
		1				QRAKALHIPFSVDEIVGMPVDS
1						FNSMLSRYYLTDLQVSLIRDIRR
						RGKNKVAAQNCRKRKLDIILNL
						EDDVCNLQAKKETLKREQAQC
	İ					NKAINIMKOKLHDLYHDIFSRL
	1					RDDQGRPVNPNHYALQCTHDG
						SILIVPKELVASGHKKETOKGK
15209	45577	Α	15299	2	342	RRRFRVAAAAVGTA*AVVLSA
				1		RPSETWETRGRR*\RICPSSWP*
				1		GI*VNNG*K*RRHSVCQCSLWF
1		l			1	PARCIKPYHSMARTQPGTRNKE
						NGPAGPTALDNVASSDDTGRH
						RPQT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15210	45578	A	15300	710	1831	IRMKSKEIIARCIKPYHSMARTO
15210	45576	ľ`	13300	1	1051	PGTRNKENGPAGPTALDNVASS
	1		İ			DDTGRHRPQTTQLAPGFAHPLQ
						LASFRRMVLFLSGEGRSRGGPE
						LOFPASCRRGEGSPGVRESGSG
	ł	l			ł	GIAATSTPNYPPNQDSKEHDIR
	l					GAEHQKQEQPAKPPHTARSAY
		l				
	Ì	l				PPQKSSYPANAKATRHSPETAA
		l				AKEARAPAAAQPQRHQPNPSP
						APHTRPATAATRQPERRVPSPT
						HRHPAATRLSPRRQSPSPRPHH
						DRRGFPRLAETLQHPMCPLPLV
		1		l		ASAGHHRHHRLLLLLAPQAPER
1	ŀ	1				EASDEI\VFSGRSRSRGCPTEFQE
1		1				SAMCFPNPGLPDSCGESQAVTI
1		1				LILRKFQKVIWVIEVPLDYKKG
	10000	١.	1.6001	77	1041	SWEYFSRMETIIMPFENIQSE
15211	45579	Α	15301	l''	1041	IPSCVVSGIAAAF/HPGLASDAP
				1		ARASSWWTHVEMGPPDPILGV
l				1		TEGFKRDTNSKKMNLGVSAYR
		1				DHNGKPYSIHKAEAQIAAKNLD
						EDYLSIGRLAEFCKASAEVALG
				ı		ENSEVSKSGRFVTVQTISGTSAL
		l]			RIRASFLQRFLKFSQDVFLPKPT
			1	1		WGSRTPIFRDTGMQLQGYRYY
						DRKTCGFDFTDTVEDISKTPEQ
1						SLLLLHACSHNPTGVDPGPKQQ
				1		KEIATVVKKT/RNLFALFDMAY
1			ł			QGFASGNGNKDAWAVRHSAN\
1						IRHCRCQSYTKNMVLHSEGVA
						GFTMVCKDADEAKRVESHLKI
15212	45580	A	15302	1	1263	LMCPMYSNPPLNGTRLLLPF
15212	45581	A	15302	i	1146	MTGRQSLSPVLSLSSLDSMSFT
13213	43381	^	13303	1	1140	TRSIFSTNYLFLGSVQVSNYGA
1				1		RPASCMASVYAGAGGSGSRISV
l				1		SRSTSFRGGMGSGDLAAGMAG
i				l		
				1		GLAGMGGIQNKKETMQSLNDR
				İ		LASYLDRVRSLETKNRKLESKI
						REHLEKKGLQVRDWSHYLKTI
1	l	1			1	EDLRVQIFANTVDNAGIVLQID
	l	1	I		İ	NACLAADDFRVKSLMTNVTWL
1		1				PLETEMEALKEELLFMKKNHEE
1		ŀ	I		1	EVKGLQAQIASSELTVEIEKSTT
		ŀ	1			VVTTQSTKVGDAEMTLTELRR
			1			TVQSLKMDLDSMRNLKASLEN
		1				SLREVEARYTLQMEQLNRILLH
		1				LESEMAQSRAEGQHQIGEYEAL
		1				LNIKVKLEAESATYHRLLDDDD
		1	1			FNLGDALDNSNSMQTI/QKTTT
				L		RRKVVSETNDTKVL

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop endon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15214	45582	Α	15304	184	370	
15215	45583	Α	15305	40	1411	RGDPRVRRRTQSLSPVLSLSPDS
						MSFTTRSTFSTNYRTLGSVQAP
	i					SYGARPVSSAASVYAGAGGSG
					İ	SRISVSRSTSFRGGMGSGGLAT
						GIAGGLAGMGGIQNEKETMQS
	l				İ	LNDRLASYLDRVRSLETENRRL
	Ì					ESKNPEHLEKKGPQVRDWSHY
						FKIIEDLRAQIFANTVEQCPTTV
						L/LRIDSA\RLA\ADDFRVKYE\T
ĺ						EAGPCAQSVENNIHGLCKVIDD
						TNVTRLQLETEIEALKEELLFM
						KKNHEEEVKGL*AQIASSELTV
						EVDAPKSQDLAEIMADIRAQYD
						ELARKNREELDKYWSQQIEEST
						TVVTTQSAEVGAAETTLTELRR
						TVQSLEIDLDSMRNLKASLENS
						LREVEARYALQMEQLNGILLHL
				İ		ESELAQTRAEGQRQAQEYEAL
		1			İ	LNIKVKLEAEIATYRRLLEDGE
						DFNLGDALDSSNSMQTIQKTTT
						RRIVDGKVVSETNDTKVLRH
15216	45584	A	15306	392	1449	GPWEMLMTIKSHPDAERQPQR
				i	i	VKKLTQVFWSSASLRER/GKTY
					!	LFWPKLKKAAEIPAAYREALDE
						ALGISGGGRDVGERVYKGPSM
						MANPQKPVLEALSSNKYLHILC
						AKHCDRHEGYKDEIKVKEASS
	ŀ					RVNGCVEAMLSAPLPLDRRNE
						HVESKGKPFTMGLSPRVLSASR
						RDGLRERKGSFFRTVVLHDTKT
				l		DSNSDTETNSSTPPRTLLEMQIL
	ĺ			1		SPTWTYEISSSRTLGCGPSVFQQ
						RASYSPSVSGPRLGCPLMDGGR
	l					MRCPPGNPESCVSEANLIPISVP
		l				VAPHAHQHLVLSVFWKITIVIG
						GKTCVAQNHSNEAKMKQRPEI
	l				l	NHASPPGAFAWSQSGEPIASAM
		L	L		l	TVPA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \-possible nucleotide insertion)
15217	45585	A	15307	I	785	MWDWGQDSSLLRGAGGPCLQL WFGRLQHLGGGLREHRDAWW HSRGLGSCSGMQGAATPTQNG RGSCLSPAPASSMEHAAPAVPP CCSWCDSSSRSWPATATTETG PRTVPSAQESAVGLLTASLLKG DSGSTGPCPLLMSLAAPGTQQC SLGPCCTFERKRSSVSPAAPGTA APELAAALMLMPRKNQIAVYE LLPKQCWWAKEGWHLPRHPE LADKNVPNLHIMKAMQISLKSQ GHREEQFAWRHFYWHLTREGI QSIQ
15218	45586	Α	15308	1	392	
15219	45587	A	15309	1	716	
15220	45588	A	15310	150	372	RRSNEWGVPVPVPARTGGG/HG QKTPPSPPLPPSSPPPSPSPPPPPA VAPAKIPPPLPLPPLPPREAPPAP GPALP
15221	45589	A	15311	218	740	KSCPETVSRRKNYGSIKKHQOD KFNPVCLKMTTKTEFVLCGWV PCRCL/REGEETVTTLDYSHCS 'LEQVPKEIFTFFKTLEELYLDA NQIEELPKQLFNCQSLHKLSLPD NDLTTLPASIANLINLRELDVSK NGIQEFPENIKNCKVLTIVEASV NGIQEFPENIKNCKVLTIVEASV NPISKLPDGFSQLLNLTQ
15222	45590	Α	15312	96	297	
15223	45591	A	15313	469	5174	KMTSLAQQLQRLALPQSDASLL SRDEVASLLPDFKEATIORDT AFAIGCTGLEELLGIDPSFEQFE APLFSQLAKTLERSVOTKAVNK QLDENISLFLIHLSPYFLLKPAQ KCLEWLIHRFHIHLYNQDSLIA CVLPYHETRIFVRVIQLLKINNS KIRWFWLLVYQSOVPLAKGT LITHCYKDLGFMDFICSLVTKS VKVFAEYPGSSAQLRVLLAFYA STIVSALIVAAEDVSDNIIAKLFP YQKGLKSSL
15224	45592	A	15314	267	495	GCLL**CLCIRL/SKNRAERALR PYSDVEALRLSSDFPFRQRPKG GSGWSLLSVEINQYYGPRVMIG RSNSEVVLVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15225	45593	Α	15315	3	3027	AMEQAGTRPAATEHPRLRRPM
						PWLLLLPLLLLLLLLPGPAASQ
				l		LRYSVPEEQAPGALVGNVARA
						LGLELRRLGPGCLRINHLGAPSF
				l		RYLELDLTSGALFVNERIDREA
				l		LCEQRPRCLLSLEVLAHNPVAV
						SAVEVEILDINDNSPRFPRPNYQ
						LQVSESVAPGARFHIESAQDPD
						VGANSVQTYELSPSEHFELDLK
						PLQENSKVLELVLRKGLDREQA
						ALHHLVLTAVDGGIPARSGTAQ
						ISVRVLDTNDNS
15226	45594	Α	15316	1	775	MPCVNGKYNSTIQAGLQMAQI
						LQDEGPYEVGIDEETGSQKGDG
						IIRSSRKLLFTMALFLETNLLKD
						PLSRERRWRLDELDSGGREKG
						VAVFLRSAGIRQVQVAMEQAG
						TRP/AATEHPRLRRPMPWLLL/L
						PLLLLLLLLPGPAA/SQLRYSV
		l				PEEQAPGAL/VGNVARALGLEL
		l	1			RRLGPG/CLRINHLGAPSPRYLE/
	1	l		l		LDLTSGALFVNERIDREACLGS
	İ	l	İ			AGANPVAVSAVEVEILDIN/DNS
						PRFPRPNYQLQVTARCWWTSW
15227	45595	A	15317	2	165	
15228	45596	A	15318	1239	2251	TCSSRARRPRRGRTVSQVRGPA
l						PWAPRRARLQQRGSLRGAARS
1		l				PASPTSA/PSYSRASASAGPAAL
						QPQGAVTPGPSANLAAPGGTL
i						VPAEGPRMPPVTAQSASSHPAP
1			}			SASQPSSPAAASRGSLSVPQEVT
1		l				ATLLASASSPTLDATRDAPATP
1						RIAAPPGSATSSRGLGSQAQGSS
		l				LGTKESISDSFGLGSPAVTQRTT
			1			APPDSAALSQDTRPEAAGSWPG
						PQTGACSVGSGTDRRDTAAAT
						EAQHLSSESKEKTSAQKSGNCF
	l	1	1			GILKADFTISTLMDPEEMKDQF
						LRQVRLPSTDALKKQSISNYRP
		l	1			PGKGKPQQYLGDSAWNMVSA
		<u> </u>				GTIEGFYKE
15229	45597	A_	15319	46	2859	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	describit, v-possible nucleotide insertion)
15230	45598	A	15320	240	924	I INGEOKCILCHIVYSSKKRH\SHE
		į.				CRRCGPTEVSLSAYAKHISGOL
						HKDNVDAQEREDDGKGEEEEE
						DYFDKELIQLIKQRKEQSRQEPS
				ŀ		SSSQEVNSDDSSSATFSVSVFYV
						WLKTILLLTWPKEVKRLDTPGL
				l		ELWLVPNMAELSEPEGPVDWK
				ł		ERCVALESQLMKFRVQASKIRE
			İ			LLAEKLHHNERLEYRDQRKDS
i		1				ASFQRKKQFTYKASGIEMASDL
		L				SKAPLEIRG
15231	45599	Α	15321	3	755	LRLPRRSTRTRRGIWVRTGAAM
						GKVNVAKLRYMSRDDFRVLTA
						VEMGMKNHEIVPGSLIASIASL
					1	KHGGCNKVLRELVKHKLIAWE
						RTKTVQGYRLTNAGYDYLALK
		1				TLSSRQVVECVG\NQMGVGK\E
1						SYIYIVANEEGQQFALKLHRLR
1						RTSFRNLKNKRDYHIHRHNVS
				ł		WLYLSRLSGMKEFCYMKQFCD
				1		EQGSQLRIHMMGGQMWAVPIC
				1		QPRDLFVWFVLRAEACAGCGA
15232	45600	A	15322	1	1028	TSQQRGALGVTSTCYRD MNKTITTQEEVLPLQADVVQRE
13232	43600	^	13322	1'	1028	EELNSLKOKLA AALLAEOEPOP
						ERLVSELQL\PRKAAVFQDEILR
				ł		SSRQLVLPELGVHRHVLVVCC
		İ		1		GGLRCPLAGGLAAASVGRLGL
	Ì			1		VDCDVVEMSNLAYQVLHGEAL
	ļ	İ		1		AAQAKAASAAASLRGLNLAPP
				1		PAETVTNCADGGMLRVLTGVL
	1					GCLQPLKVLKIAASLGPSYSGS
				Į.	l	LLLFDALRGHFCCIRLRSCRLDC
				İ		AACGKWPTVTHLLDYEAFCGS
		1		l		SATDKCRSLQLLGPEDRVSVTD
				į.		YKRLLDSGAPHLLLDIRPKMEL
		i		1	1	PRKAAVFHDEILRSSRQLVLPEL
				1	1	GVLRHLLVVGCGGLRCPLAQY
				ł		LAAAIVGRLGLVAYDVV
15233	45601	Α	15323	92	778	QGTRRWSLGTCSLMTRTQFEV
						CTKVCVQESLQRPGEELKTRCP
				1		SVGAGYVNYGTPAHERI/GAAV
						KTGREADKPEVTKQTQFSTRK
				1	1	DE/TSLFGVPLQ*LLARHRAPP*
		1		1	1	RRT/RRLLEVARTPMLPAQASR
				1		SELAARQDSAWASPAAPASPN/
1		1		1		RSVLRRCLLSARPPLCGPL/LRV
1				1	1	LPSWRRQQIRENSAED*LHSEY
1			1	1	1	RANRIRRRRGHPEN/WTKS*LE
		L.				NPENLTRRSSSAPAPC
15234	45602	Α	15324	142	500	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15235	45603	Α	15325	3	343	
15236	45604	Α	15326	1164	1648	PLMTSAPWPSCCCWRSLSSGRP
		1				AGRQRQLPTSITTANQPAPSRR
		1		Į.		KKSTSRKKQRNLPAVSPAMVAI
		1				V\RRSCDLYRWSVLHYASQERR
		1				VRDAAKPESD\RKRTTTKTRR/H
		1				WRYIKELESRQPGVADMRQQP
		1				TQLVEVPWNEQTPEQRQQTLQ
15037	45505	ļ	16227	20	417	RQHGRETRCSV
15237	45605	A	15327	28	440	
15238	45606 45607	A B	15328	1	440	
15240	45608	A	15339	22	584	IRGRVDPPSAVEPPPKMQIFVKT
13240	43008	Α.	13330	22	364	LTG\KT\ITLEVEPLGYG*KNVK
						AOASRDK\EGI\PPDOORL\IFAG
		1				KQ\LEDGRTLS\DYNIQK\ESTL/
						HIFVLRLRGGAKKRKKKSYTT\
l						PKK\IKHKRKKG*SLAV\LKYYK
		1				\VDE\NGKIS\RLRRECPS\DECGC
-						WGVYGOVTFDRHYCG\KCCLT
						LLFQQTRMTSNCMS
15241	45609	la	15331	3	954	LDARRHWEGTPGFTHTSPDAW
					,	ADAWGKMAGEKVEKPDTKEK
				1	İ	KPEAKK\VDAGGKVKKGNLKA
	1					KKPKKGKP\HCSRNPVLLRGIG
						RYSRSAMYSRKAMYK\RKYSA
						AKSKVEKKKKEKVLATVTKPV
				ı		GGDKNGGTRVVKLRKMPRYY
				1		PTEDVPRKLLSHGKK\PFSQHV
						RKLRASITPGTILIILTGRHRGKR
		1		ŀ		VVFLKQLASGLLLVTGPLVLNR
		1				\VPLRRTHQKFVI\ATFTKIDISN\
		1				VKIPKHLTDAYFKKKKLRKPRH
						QEGEIFDTEKEKYEITEQRKIDQ
						KAVDSQILPKIKAIPQLQGNLRS
		ـــــــــــــــــــــــــــــــــــــ				VFALTNGIYPHQLVF
15242	45610	Α	15332	670	1008	WCILGVTPSLMSRTALFLSRHT
l		1		1	1	FFA/RSHSAASKL/EKKKKEKVL
		1		1	1	ATVTKPVGGDKNGGTRVVKLR
		ı		1		KMPRYYPTEDVPRKLLSHGKK
l		1		1		PFSQHVRKLRPSITPG\TILIILTG\
	45511	١	1.5000	 	700	RHRG
15243	45611	Α	15333	1	708	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15244	45612	Α	15334	Ĭ	1754	MRNEISDLTEELHQKEITIATVT
						KKAALLEKQLKMELEIKEKML
						AKQKPWACLVNAMPAIQVVFN
						LGRFRKCQDPHKAQEWVYPGY
			l			SGSDCEVFPEIMTSKNREEAMP
						EWKLWKKVVLSGATDAPSWD
1		ĺ				GPTFTDLNVFAGNMGLSDCTLS
l						LGKIWKIPGHFLHTSGPVSFHT
						KGLLNTPTKERWVGGFGSEGR
						GVPEPGGCFGVLAVPRRTPPSA
						AAMTGGERLGRAWAGDRAAA
						SRSAPVPGTFGMMGWRPVAFG
						KNAKAYRGFGGIVTSLTPSCDK
				ŀ		VTLLFVRILPGSGQSRDQSSPEK
						LTDTKSTSNKAYGFPECLRRPE
						STHIFFYDHQLGSEQPRGTKGT PLKWVSQKTLERETRQCQALVI
						WTDCDREGENIGFEIIHVCKAV
						KPNLOVLRARFSEITPHAVRTA
		1				CENLTEPDQRVSDAVDVRQEL
						DLRIGAAFTRFQTLRLQRIFPEV
			l			LAEQLISYGSCQFPTLGFVVERF
						KAIOAFVPEIFHRIKVTHDHKD
				1		GIVEFNWKR/LSTL*PHGLPSSL
1						SVVCGVRSLSCRSVGFCWRSTP
		1				DPVCLCITSGGCRTASIAEOOKL
						LPDPSYRSFIPEGQPPI
15245	45613	Α	15335	3	125	
15246	45614	A_	15336	764	1005	
15247	45615	Α	15337	1	3533	MELVNCVLMSTLGNTCCPMAI
1						AMFRSSSSLPRPGHSLTGSPLRV
						NLIHIKFPLNDFHQNVVKIKNK
						DPPLGAAIGKESCPIADNELHSL
						ERRPLSPKLVQALGVAGLSRPG
						PPHTGCAIWLLLITDHKLLSEIL
						YDLLHQGLDILADPPDEGGIPR
		1	l			GKNLVELLQLPGLEEDKAFQKE
		1	l			IGLKILVFKAYSTLSSDGPPLFS
1		l	ŀ	l		ALRSLQGVQSSKVGESREEKSQ
1						TEEEDPGNPWVPQSLGASVSSS
1						GKWGCLRTKP

NO:	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
15248 45616 A 15338 1 2800 SISGPGQEAVPLRPKAEPPEGS MVWDRQTKMEYEWKPDEQC QQLQLLCESQSPOTTIGRTVQ KLEQLNQYPDFNNYLIFVLTK KSEDPTRISLSGLIKNNVKAI FQNFPNGYTDFKSECLNNIGT SPLIRATVGILITTIASKGELQN WPDLPKLCSLLDSEDYNLTE AFGALQKICEDSAEILDSDVLT RPLNIMPKFLQFKHSSPKIRS AVACVNGFIISRTQALMILHIDS TENLFALA 15339 8 450 TWSWSPQAPSWARPSSQARA SLLVARQQLCQGSRRRSGVGS VDAGAARTALRTLGTRPRIKT CGIGSHETKAAASGQDQGKG RGRGSLGGRELGANEPLYQW AHTLTVLFILICTLGYVTLLEE PQDTAYNTKSEIAVPHLS 15250 45618 A 15340 246 559 MPYAVMAESSTAFKKLDQG NQELEAPGGIATPPVYGQLLA YLLHN/GQRIWARPSFKKLDQG NQELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDQG NQELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDQG NGELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDQG NGELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDGG NGELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDGG NGELEAPGGIATPPVYGQLA YLLHN/GQRIWARPSFKKLDGG NGELEAPGGIATPFVYGQLA YLLHN/GQRIWARPSFKKLDGG NGELEAPGGIATPFVYGQLA YLLHN/GQRIWARPSFKKLDGG NGELEAPGGIATPFVYGQLA ALYLHHNDMNAARVL*KRIPP IKSANSELGGIWSVGQRIWAR MPYAVMAESAVSFKKLDGG NGELEAPGGIATPKVYGQLI ALYLHHNDMNAARVL*KRIPP IKSANSELGGIWSVGQRIWAR GFWCRGPWDYFPNFKISLY PAPAVPPIPPREQULARLTDVYAFLEN NGENGWGWADSTTRINGSCI GKPVCRGPWDYPPNFKISLY PAPAVPPIPPREQULARLTDVYAFLEN NGENGWGWGWADSTTRINGSCI GKPVCRGPWDYFPNFKISLY PAPAVPPIPPREQULARLTDVYAFLEN NGENGWGWGWGWGWGWGWGWGWGWGWGWGWGWGWGWGWGW						codon for last amino acid	
15248 45616 A 15338 1 2800 SISGPGQEAVPLRPKAEPPEGG				09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
Mywdrqtkmeyewdedord					sequence		
Mywdrqtkmeyewdedord	15240	10010	_	licano		2000	DISCOURT AND DRY A EDDESCO
QQILOLLKESQSPOTTIQRTVC KLEQLNQYPDENYVLIFVLTK KSEDEPTRSLSGLILKNNVKA FQNPPNGYTDFIKSECLNIGG SPLIRATVGILITITASKGELQN WPDLIPKLCSLLDSEDYNTCE AFGALQKICEDSA BILDSDVLT RPLINIMPKELQFFKHSSPKIRS AVACVNQFIISRTQALMI.HID TENLFALA A	15248	45616	А	15338	11	2800	
KLEQLNQYPDENNYLIFVLTK KSEDEPTRSLSGLIKNNVKAI FQNPPNGYTDFIKSECLNINGT SPLIRATVGILITITIASKGELQN WPPDLPKLCSLLDSEDYNTE AFGALQKICEDSAEILDSDVLT RPLNIMIPKELQFFKHSPKIRS A VACVNQFIISRTQALMLHIDE TENLEALA TENLEALA TENLEALA TENLEALA TENLEALA STANGOLCOGSRRRSGVGS VDAGARTALRTGTRPRIRE CGLGSHETKAAASQDGQRG RGRGSLGGRELGANEPLVQW AHTLTVLFILTCTLGYVTLEE PQDTAYNTKSEIAVYHLS PQDTAYNTKSEIAVYHLS PQDTAYNTKSEIAVYHLS TAHOWSETVQPIMEALRDATRI A 15340 246 559 MPVAVMAESAFSKKLDQG QELEAPGGIATPPVYQGLIA YLLINOQRIWQRDPFQIYTIT AHQWSETVQPIMEALRDATRI RAFALVSQAYTSIIADDFA RSGTQGSSAGTAARALNVIAGAI TAHOWSETVQPIMEALRDATRI TS251 45619 A 15341 129 915 RSGTQGSSAGTAAVTVQRRGPAK MPVAVMAESAVSFKKLDQG ENQELEAPGGATPFVYQQLIA ALVLFHNDMNARAVLYRIPP IRSANSELGGIWSVQGRIWQB DFFWGGYTTINGHQWSETVQ IMMEALBATRETAPLEWVSQ, YTSIIAR*FCQPLLGLPVIEAA KGILEQGWQAADSTTRNGSCI GKPVCRGPWDYPPNFFISLS, EPAPVPPIPPREQLARLTDV AFLEN TS252 45620 A 15342 1 529 MATLTISRAQTEDEADYVCHR KLVKEGLDERTHKAYLSSGK GCEFHMVKROSPLGPPILGSW QSALIQPPSVSGSPGQSVTISCT GTSSDVGSYDYVSWYQQHRG VPKPMIYNVTQPSRVPDRFSS SKSGWTTSAMTISCLQAEDEAL YYCCSYAGSPYVFVGGTGKL TS253 45621 A 15343 399 737 WISGGNSCATTIJSDAGLSTTS SIGSATGARQAESASSOSGG LGRHCGASGKCAESLRREPVC ATTQWAPGWVVCRPASDSSH VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSSWSPAEVYPSLEIPER VIGGRSWSPAEVYPS							
					1		
FQNFPNGVTDFIKSECLNNIGT SPLIRATVGILITITIASKGELQN WPDLPKLCSLLDSEDYNTG AFGALQKICEDSAEILDSDVLT RPLNIMPKFLQFFKHSSPKIS AVACVNGFIISRTQALMLHIDG TENLFALA TWSWSPQAFWARPSSQARA SLLVARQQLCQGSRRRSGVGS VDAGARTALRITLGTPRPLRIT CGLGSHETKAAASGQDQGGG RGRGSLGGRELGANEPLYQW AHTLIVLFILITCTLGYVTLEE PQDTAYNTKSEIAVVHLS SSEGNAFFKKLDQF VDAGARTALRITGTPRPLRIT CGLGSHETKAAASGQDQGGG RGRGSLGGRELGANEPLYQW AHTLIVLFILITCTLGYVTLEE PQDTAYNTKSEIAVVHLS SSEGNAFFKKLDQF VQELEAPGGIATPPVGQLLA YLLHN/GQRIWQRDFPGIYTT AHQWSETVQPIMEALRDATTI AHQWSETVQPIMEALRDATTI AHQWSETVQPIMEALRDATTI AHQWSETVQPIMEALRDATTI AHQWSETVQPIMEALRDATTI AHGWSETVQPIMEALRDATTI AFALVSQAYISIIADDFA SGCTQGRSAATAALNIVIAGAI EGGSWGLAVRTVQRRGPAK MPVAVMAESAVSFKKLLDQG ENQELEAPGGIATPKVYGQLL ALVLFHNDMNAARVL*KRIPP IRSANSELGGIWSVGQRIWG DFFWGYTTTINGHQWSETVQ IMMEALBDATRETAPLPWVSQ, YTSWIAR*FCQPLLGLPVIEEAV KGILEQGWQADSTTRNSCI GKPVCRGPWDYPPNFRISLY EPAPVPPIPPREQLARLTDYV AFLEN SECTION AFLEN SECTION AFLEN SECTION SECT							
SPURATYGILITIASKGELON WPDLIPKLCSLLDSEDYNTCE AFGALQKICEDSA EILDSDVIT RPLNIMIPKFLQFFKHSSPKIRS AVACVNQFIISKTQALMH.IDD							
WPDLLPKLCSLLDSEDVNTCE							
AFGALQKICEDSAEILDSDVLI							SPLIRATVGILITTIASKGELQN
RPLNIMIPKELOFEKHSSPKIRS				ŀ			WPDLLPKLCSLLDSEDYNTCEG
AVACVNĢFIISRTQALMLHIDS TENLFALA 15249 45617 A 15339 8 450 TWSWSPQAPSWARPSSQARA SLLVARQQLCQGSRRRSGVOS VDAGAARTALRILGTRPRIRE CGLGSHETKAAASGQDQGG RGRGSLGGRELGANEPLYQW AHTLIVLFILICTLGYVILLE PQDTAYNTKSEIAVYHLS PQDTAYNTKSEIAVYHLS 15250 45618 A 15340 246 559 MPVAVMAESSAFSKKLDQG NPVAVMAESSAFSKKLDQG NQELEAPGGIATPPVYQGLLA YLLHN/GQRIWQRDFPGIYTTI AHQWSETVQPIMEALRDATRI ARFALVSQAYISIIADDFA RSGTQGRSAATAALNIVIAGAI 15251 45619 A 15341 129 915 RSGTQGRSAATAALNIVIAGAI RAFALVSQAYISIIADDFA RSGTQGRSAATAALNIVIAGAI RAFALVSQAYISIIADDFA RSGTQGRSAATAALNIVIAGAI ALVLFHNDMNAARVL*KRIPP IKSANSELGGIWSVGQRIWG DFFWG/IYTTINGHQWSETVQ IMMEALBDATRETAPLPWVSQ, YTSWIAR*FCQPLLGLPVIEEAV KGILEQGWQADSTTRNGSCI GKPVCRGPWDYPPNFRISI, YEPAPVPIPPNEQULARLTDVY AFLEN S29 MATLTISRAQTEDEADYVCHF KLVKEGLDERTHKAYLSSGK GCEPHMVKPGSPLGPDLIGSW QSALIQPSVSSGFQGSVTISCT GTSSDVGSYDYSWYQQHEG VPKPMIYNVTQPSRVPDRFS SKSGWTTASMTISCLQAEDEAL YYCCSYAGSPYVSWYQQHEG 15253 45621 A 15343 399 737 WISGGNSCATTIJSDAGLSTTS SIGSATGARQAESASSQSGG LGRHCASGKCAESLREPVC ATNQWAPGWVVCRRASDSSH VIGLGRSAWSPAEVTPSLIPCR							AFGALQKICEDSAEILDSDVLD
TENLEALA							RPLNIMIPKFLQFFKHSSPKIRSH
15249 45617 A 15339 8 450	ĺ						AVACVNQFIISRTQALMLHIDSF
SLLVARGOLCQGSRRRSGVG							TENLFALA
VDAGAARTALRTLGTRPRLRF CGLGSHETKAAASGQDGQRG RGRGSLGGRELGANPEN_CVV AHTLTVLPILTCTLGYVTLLEE PQDTAYNTKSELAVYHLS 15250 45618 A 15340 246 559 MPVAVMAES/AFSFKKLLDQC NQELEAPGGIATPPVYQGLLA YLLIHN/QQRIWQRDPFGYTYII AHQWSETVQPIMEALRDATRI RAFALVSQAYISILADDFA TS251 45619 A 15341 129 915 RSGTQGRSAATAALNVIAGA1 EQQSWGLAVRIVQRGPAK MPVAVMAESAVSFKKLLDQC ENQELEAPGGIATPKVYQGLL ALVLFHNDMINARVI_KRIPP IKSANSELGGIWSVGQRIWQR DFPWG/IYTIINGHQWSETIVQ IMBEALRDATREAPLPWNSQ, YTSIIAR*PCQPLLGLPVIEFAX KGILEGGWQADASTTRNGSG GKPVCRGPWDVFPFNKFISLY: EPPAPVPIPPREQLARLIDIVA AFLEN 15252 45620 A 15342 1 529 MATLITISRAQTEDEADYYCHE KLVKEGLDERTHKAYLSSSG GGFFHMVKPGSPLGPDILGSW QSALIQPSVSGSPGQSVTISCI GTSSDVGSYDYVSWYQQHEG VPKPMIYNVTIQPSRVPDRFS SKSGNTASMTISGLQAEDEAL YYCCSYAGSYPVVFGGGTKLI 15253 45621 A 15343 399 737 WISGGNSCATTIJSDAGLSTITS SIGSATGARQAESASSQSGG LGRHCASGKCAESLRREPVC ATNQWAPGWVYGRPASDSSH VIGLGRSAWSPAEVTPSLIPCR	15249	45617	A	15339	8	450	TWSWSPQAPSWARPSSQARAV
CGLGSHETKAAASGODGORG RGRGSLGGRELGANEPLYQW AHTLTVLFILITCTLGYVTLEE PQDTAYNTKSEIAVVHLS	1						SLLVARGQLCQGSRRRSGVGSR
RGRGSLGGRELGANEPLYQW AHTLTVLFILTCTLGYVTLEE PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS NELEAPGGIATFPYVGQLLA YLLHN/GQRIWQRDFFGIYTT AHGWSETVQFMERALAT YLLHN/GQRIWQRDFFGIYTT AHGWSETVQFMERALAT AFLVSQAYTSIIADDFA FALVSQAYTSIIADDFA FALVSQAYTSIIADDFA FEGSWGLAVRTVQRRGPRAS MPVAVMAESAVSFKKLLDQC ENQELEAPGGIATFEKYVGQLC ENQELEAPGGIATFEKYVGQLC ALYLFHNDMNARYL-KRIPP IKSANSELGGIWSVGQRIWQR DFPWG/YTTINGHOWSETVQ TSIMIAR FEQPLLGLPVEEAN KGILEGGWQAADSTTRNGSC GKPVCRGPWDVFPNKFISLY: EPAPVVPPIPNEQQLARLTDV AFLEN FEGSWGLAVTKHAYLSSGA GCFHMVKPGSPLGPDILGSW QSALIQPSVSGSFGQSVTISCT GTSSDVGSYDYVSWYQQHEG VPKPMIYNVTNGPSRVPDRFS KSGNTASMTISCLQAEDEAL YYCCSYAGSYPVVFGGGTKLT YYCCSYAGSYPVVFGGGTKLT SIGSATGARQAESASSOSGG LGRHCASGKCAFSLRREPVC ATNOWAPGWVVGRPASDSSH VIGLGRSAWSPAEVTSELDEGE	1						VDAGAARTALRTLGTRPRLRRP
RGRGSLGGRELGANEPLYQW AHTLTVLFILTCTLGYVTLEE PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS PQDTAYNTKSEIAVVHLS NELEAPGGIATFPYVGQLLA YLLHN/GQRIWQRDFFGIYTT AHGWSETVQFMERALAT YLLHN/GQRIWQRDFFGIYTT AHGWSETVQFMERALAT AFLVSQAYTSIIADDFA FALVSQAYTSIIADDFA FALVSQAYTSIIADDFA FEGSWGLAVRTVQRRGPRAS MPVAVMAESAVSFKKLLDQC ENQELEAPGGIATFEKYVGQLC ENQELEAPGGIATFEKYVGQLC ALYLFHNDMNARYL-KRIPP IKSANSELGGIWSVGQRIWQR DFPWG/YTTINGHOWSETVQ TSIMIAR FEQPLLGLPVEEAN KGILEGGWQAADSTTRNGSC GKPVCRGPWDVFPNKFISLY: EPAPVVPPIPNEQQLARLTDV AFLEN FEGSWGLAVTKHAYLSSGA GCFHMVKPGSPLGPDILGSW QSALIQPSVSGSFGQSVTISCT GTSSDVGSYDYVSWYQQHEG VPKPMIYNVTNGPSRVPDRFS KSGNTASMTISCLQAEDEAL YYCCSYAGSYPVVFGGGTKLT YYCCSYAGSYPVVFGGGTKLT SIGSATGARQAESASSOSGG LGRHCASGKCAFSLRREPVC ATNOWAPGWVVGRPASDSSH VIGLGRSAWSPAEVTSELDEGE							CGLGSHETKAAASGQDGQRGS
PQDTANTKSEIAVYHLS	l						RGRGSLGGRELGANEPLYQ/WR
15250 45618 A 15340 246 559 MPYAVMAES/AFSFKKLLDQC NQELEAPGGIATPPVYGQLLA YLLIH/NQRIWQRDFPGIYTIT AHQWSETVQPIMEALRADATRI RAFALVSQAYYSIIADDFA 15251 45619 A 15341 129 915 RSGTQGRSAATAALNVIAGAT EQQWGLAVRTVQRRGRPAK MPYAVMAESAVSFKKLLDQC ENQELEAPGGIATPKVYGQLL ALVLFHNDMINARVL*KRIPP IKSANSELGGIWSVGQRIWQR DFPWG/IYTTINGHQWSETNVQ IMEALRDATRETAPLPWVSQ, YTSIIAR*PCQPLLGLPVIEFAN KGILEQGWQADASTTRINGSC GKPVCRGPWDVFPFNKFISLY: EPYAPVVPIPPNEQQLARLTDTV AFLEN 15252 45620 A 15342 1 529 MATLITISRAQTEDEADYYCHE KLVKEGLDERTHKAYLSSSG GCEFHMVKPGSPLGPDILGSW QSALIQPSVSSSPGQSVTISCT GTSSDVGSYDYVSWYQQHEG VPKPMIYNVTQPSRVPDRFS SKSGNTASHTISLQAGEDEAL YYCCSYAGSYPVVFGGFTKLT SIGSANTGARQAESASSQSGG LGRHCASGKCAPSLRREPVG ATNQWAPGWVVGRPASDSSH VIGLGRSAWSPAEVTSELDEGE							AHTLTVLFILTCTLGYVTLLEET
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QHCKIHTREKPYECIKCCKAFS THSSYVQHLKIHTGEKIHECNQ CGKAFSHSSNLIHHQRIHSGEKP YKCKECGKAFNRQSNLIQHQRI HSGEKPYDCKECGKAFSTOGHL IQHQRIHTGEKPYECNECAKSFS LNRTLTVHQRIHTGEKPYECNECAKSFS LNRTLTVHQRIHTGEKPYECNECAKSFS CGKSFSQCSQVIQHKRIHTGEK PYICNECGKSFGARLSLIQHQRI HTGEKPYGCREKPYECSECGK AFSQSFNLIHHQRTHNGEKSFY CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLQ HQRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEKSFYEC KAFSQSAHLIHRQRITGENFYP ECSECGKAFNVCSSLIQHIRIHT GKPYECSDCG KAFSQSAHLIHRQRITGENFYP ECSECGKAFNVCSSLIQHIRIHT GKPYECSDCGKAFSQHSHQH HQRIHTGEKPYECSDCG	ł													
THSS YOÇHLKIHTGEKHECNO CGKAFSHSNLIHHQRIHSGEKP YKCKECGKAFNGSNLIQHQRI HGEKPYDCKECGKAFSTQLFL IQHQRIHTGEKPYENCEAKS LINTLITHQRIHTGEKPYENCEAKS LINTLITHQRIHTGEKPYENCE CGKSFGQCSQVIQHKRIHTGEK PYICNEGGKSFGARLSLIQHQRI HTGEKPYGCEKPYECSECGK AFSQSFNLIHINGTHNGEKSYE CNECDKAFSLLSSLVQHQRIHN GDKYYECHKCGKAFSQGSHLIQ HQRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSANLIHHQRIHTGERY EGSEGGKAFNVCSSLIQHRRIHT GGKYYECSDCGKAFSQHSQHIQ HQRSHIGEKPYECSDCG KAFSQSANLIHQRIHTGERYPYCSDCG KAFSQSANLIHQRIHTGEKPYMCNECEKSFSA	ł		1			i								
CGKAFSHSSNLIHHORIHSGEKP YKCKECGKAFNRQSNLIQHQRI HGGEKPYDCKECGKAFSTQLFL IQHQRIHTGEKPYECNECAKSFS LNRTLTVHQRIHTGEKPYCNEC CGKSFSQCSQVIQHKRIHTGEK PYICNECGKSFGARLSLIQHQRI HTGEKPYGCREKPYECSEG AFSQSFNLIHHQRTHNGEKSYE CNECDKAFSULSSLVQHQRIH GDKPYECHKCGKAFSQGSHLIQH HQRSHIGEKPYECSECG ISTLIKHERTHNGEKPYECSDCG KAFSQSALIHHQRIHTGENPY ECSECGKAFNVCSSLIQHRIHIT GEKPYCSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA														
VKCKECKAFNRQSNLIQHQRI HSGEKPYDCKECGKAFSTQLFL IQHQRIHTGEKPYECNECAKSFS LNRTLTVHQRIHTGEKPYRCNE CGKSFSQCSQVIQHKRIHTGEK PYICNECGKSFGARLSLIQHQRI HTGEKPYGCREKPYECSECGK AFSQSPNLIHHQFRIHTGEKPY CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLIQ HQRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSAHLIHRQRHTGEKPY ECSECGKAFNVCSSLIQHHRIHT GEKPYECNDCGKFFQHQHRIHTGEKPYENDCH			1											
HSGERPYDCKECĞKAPSTOLFL IQHQRIHTGEKPYECNECAKSPS LNRTLTVHQRIHTGEKPYECNEC CGKSFSQCSQVIQHKRIHTGEK PYICNECKSFSGARLSLIQHQRI HTGEKPYECSECGK AFSQSPNLIHHQFSKYE CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCĞKAFSQCSHLQ HQRSHIGEKPYECNECGKTFQ ISTLIKHERTHNGEKSYE CNECDKAFSULSILVQHQRIHN GDKPYECHKCĞKAFSQCSHLQ HQRSHIGEKPYECSDCG KAFSQSAHLIHQRIHTGEKPYECSDCG KAFSQSAHLIHQRIHTGEKPYECSDCG KAFSQSAHLIHQRIHTGEKPYECSDCG HQRIHTGEKPYMCSSLIQHHRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCSSLIQHHRIHT						1								
IQHQRIHTGEKPYECNECAKSPS LNRTLTVHQRIHTGEKPYRCNE CGKSFSQCSQVIQHKRIHTGEKP PYICNECGKSFGARLSLIQHQRI HTGEKPYGCREKPYECSECGK AFSQSFNLIHHGRTHNGEKSYE CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLQ HQRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSAHLIHRGRITGENPY ECSECGKAFNVCSSLIQHHRIHT GEKPYECSGCGKAFSQSHLGHQRITGERPY HQRHTGEKPYHCNECGKTFGQ WARNEL						1								
LNRTLTVHORIHTGERPYRCNE CGKSFSQCSQVIQHKRIHTGER PYICNECGKSFGARLSLIGHQRI HTGEKPYGCREKPYECSECGK AFSQSNLIHHQRITHGEKSY CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLIQ HQRSHIGEKPY ECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSALIHIRGRIHTGENFY ECSECGKAFNVCSSLIQHIRIHT GEKPYECDGCAFTGQ HQRHTGEKPY HCRECKFFGARTGGERPYECSDCG KAFSQSALIHHGHTGENFY ECSECGKAFNVCSSLIQHIRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA														
CGKSFSQCSQVIOHKRIHTGEK PYICNECGKSFGARLSLIQHQRI HTGEKPYGCREKPYECSECGK AFSQSFNLHIRQRTHNGEKSYE CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLQJ HQRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEK PYECSDCG KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHIRIHT GEKPYECDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA														
PYICNECÓKSFGARLSLIÓHÓRI HTGEKPYGCREKPYGCSECGK AFSQSFNLIHHÓRTINGEKSYE CNECDKAFSLLSSLVÓHÓRIHN GDKPYECHKCÓKAFSQÓSHLÓJ HÖRSHIGEKPYECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSAHLIHHÖRITTGENPY ECSECGKARNVCSSLIÓHHRIHT GEKPYECSDCGKAFSQHAHÖRITGENPY HÖRITGERPYECSDCHAFÓRIHIHTT GEKPYECSDCGKAFSQHAGÐIÐ HÖRIHTGERFYMCNECEKSFSA														
HTGEK PYGCREK PYECSECGK AFSQSFNLIHORTHNGEKSYE CNECDKAFSLLSAL VOHORIHN GDK PYECHKCGKAFSQGSHLIQ HQRSHIGEK PYECNECGKTFGQ ISTLIKHERTHNGEK PYECSDGG KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHIRIHT GEK PYECSDCGKAFSQHSQFIQ HQRIHTGEK PYMCNECEKSFSA														
AFSQSFNLIHHQRTHNGEKSYE CNECDKA FSLLSSLVQHQRIHN GDRPYECHKCGKAFSQGSHLQI HQRSHIGBERY ECNECGKTFQQ ISTLIKHERTHNGEK FYECSDCG KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHIRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA	İ		1											
CNECDKAFSLLSSLVQHQRIHN GDKPYECHKCGKAFSQGSHLQ HQRSHIGEKPY ECNECGKTFGQ ISTLIKHERTHNGEKPYECSDCG KAFSQSAHLIHRQRHTGEKPY ECSECGKAFNVCSSLIQHHRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPY			1				HTGEKPYGCREKPYECSECGK							
GDKPYECHKCGKAFSQGSHLIQ HQRSHIGEKPY ECNECGKTFGQ ISTLIKHERTHSGEKPYECSDCG KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHHRIHT GEKPYECSDCCKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA							AFSQSFNLIHHQRTHNGEKSYE							
HQRSHIGEKPY ECNECÖKTFGQ ISTLIKHERTINGEKPY ECSDCG KAFSQS AHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHIRIHT GEKPY ECSDCGKAFSQHSQFIQ HQRIHTGEKPY MCNECEKSFSA HQRIHTGEKPY MCNECEKSFSA							CNECDKAFSLLSSLVQHQRIHN							
ISTLIKHERTHNGEK PYECSDCG KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHHRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA			1											
KAFSQSAHLIHHQRIHTGENPY ECSECGKAFNVCSSLIQHIRIHT GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA	1		1		l									
ECSEČGKARNVCSSLIGHIRIHT GEKPYECSDCGKARSQHSQFIQ HQRIHTGEKPYMCNECEKSPSA	1		1				ISTLIKHERTHNGEKPYECSDCG							
GEKPYECSDCGKAFSQHSQFIQ HQRIHTGEKPYMCNECEKSFSA			1		I		KAFSQSAHLIHHQRIHTGENPY							
HQRIHTGEKPYMCNECEKSFSA	1	1	1		l		ECSECGKAFNVCSSLIQHHRIHT							
	1		1		l		GEKPYECSDCGKAFSQHSQFIQ							
			1				HQRIHTGEKPYMCNECEKSFSA							
CLSLIQHKRIHTGEKPYVCAKC			1		l		CLSLIQHKRIHTGEKPYVCAKC							

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
15259	45627	Α	15349	[3	354	EAFGGERVSSTMSGLSGPPARR GPFPLALLLIFLLGPRLVLAISF HLPINSRKCLREEIQQGPASDW RVRDLRPVWGRWRPAQ/RTSRS QILLAIFSTPKRMQPRGNLPLTT EDYDMF
15260	45628	Α	15350	283	393	YVYRRYGRFE*NDPSSALMST* GKRCQDCLDQRENKK
15261	45629	A	15351	1	351	
15262	45630	Α	15352	1	867	
15263	45631	В	15353	1	1323	
15264	45632	Α	15354	1	3669	
15265	45634	A	15355	74	208	LESYLQLPSAEGKTVSVFGETC ATPVGP/AAGPHTQ/LAQNIVTS WLTGGRFIELKTVQ/ILDRLELE KPCIDAEDECFNTEWSTEFTLL KAWDEYLKAWFALH.LEAMF QPSDSGKSFIFNMSVGYNLEGQ PLNPKNYPSGGVPRVLKSHRQD VLYGNKLSWADJHLVELFYYV EELDSSLISSFPLLRPHTHVDNT KKGSHPHMCAYTDYVNNPND RMPQREITEVS QRLTKLRETSRRLLSWQRSPSS TISMHGAEWSPPSWLLAAAGV
15267	45635	A	15357	308	735	FEEKFIKSAEDLDKVRNDGYL MFQQVP\RLRLMG*SWSRV NQRCTWAYSMTMEIFLISCVKT LSVRYAVQOLKTGVFCSLETOP
						SKMAPAKKGSKKKKGHS/AIIE VVTQENTINIQKHIHEVGFKKC APRALKEIQKFAMKEMGTPVV LIDTRLNKAVWAKRIRNVTNRI HLCLSRKCNEDKE
15268	45637	В	15358	3	991	LSSLASMSFTTCSAFTNYWSPG SVQVPSYGTQPVSHAASVYAG LGGSGSRISVSHASPMAGGLAG MGGIQNEKETMQSLRDRLASY LDRVRGLETENWKLESKIQEH

	01/0/3007					FC1/U301/08031
SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540.217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	deterion, (-possible nucleotide insertion)
15270	45638	A	15360	40	1030	RGDPRVRRRTOSLSPVLSLSPD
						MSFTTRSTFSTNYRTLGSVQAF
	1					SYGARPVSSAASVYAGAGGSC
						SRISVSRSTSFRGGMGSGGLAT
						GIAGGLAGMGGIQNEKETMOS
						LNDRLASYLDRVRSLETENRR
	1					\ESKIREHLEKKGPQVRDWSH\
						FKIIEDLRAQIFANTVDNARIVI
				l .		QIDNARLAADDFRVKYETELA
						MROS\VEN\DIHGLRNVI\DDTN
						YHTDLOLETEIEALKEELLFMK
						KNHEEEVKGLQAQIASS\GLTV
	l	l		i		EV\DAPK\SQDLAKIMA\DIRAQ
		l				Y\DELAR\KNREE\LDKYWSOO
		l				EESTTVVTTOSAEVGAAETTLT
15271	45639	A	15361	1	438	LHSDDLSLOMLRCSPVATAAS
132/1	43037	^	13301	ľ	130	TSGPGAPISSSSFSRVGSSSFQG
	i	1		l		LGGGFGGASGIGGITAVMVNO
						LLRPLNLEVDPNIQARF/VHTQ
		1		1		KEQIKTF/NNKFASFIDKVRFLE
		1				OONRMLEAKWSLLOOOKMAG
						SNMDNMFQSYINNL
15272	45640	A	15362	176	1455	EIFPLVIOKSYKVSTSGPRAFSS
13272	13040	ľ.	13302	1,,,	11433	R\SYTSGPGAPISSSSFSRVGSSS
		l				FOGGLGGGFGGASGIGGITAV
						/VQVLHTQEKEQIKTF/NNKFAS
						FIDKVRFLEQONRMLEAKWSL
						LOOOKMAOSNMDNMFOSYIN
						NLRWQLETLGQEKLKLEAELG
						NMQELVEDFKKKYHDEINKHT
		ļ				EMENEFVLIEKDVDEAYKNKV
		1				ELESHLEGLTDEINFLRQLHEE
	į.		į.			IWELQSLISDTSVVLSIDSSHSL
	1					MDNIITEVKAOYKEIANCSWA
	1	1				ADSMYQIKYEDLQMLARKHG
	İ	1				NLRCTKTDISEMNQNVSWLQA
	1	1				EIKGLKGQRASLEATITDAEKR
	1					RELAIKDANTKLLELEAALQW
	1				1	AKQDMAQQLRVYQELMNVKI
	1					ALDIKTATYKKLLEGEESWOE
	l l	1	İ		1	RMQNMSIYSKTTSGYAGGLSS
	1	1			1	AYGGLTSPP\LSYGLSS
15273	45641	В	15363	70	915	5521511 2516255
15274	45642	C	15364	62	217	
15275	45643	В	15365	1	612	
132/3	43043	ь	13303	<u> </u>	012	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15276	45644	۸	15366	3	841	STPKMPKGKAKGKKVAPAPA VYKKQBAKVVNPLEEKERPEN FGIGQDIQPKRDLTRFVKWPRY IRLORQRAILYKRLKVPPAINOF IRLORQRAILYKRLKVPPAINOF ETKQBEKQRILUARABKKAAW PKGTFPTKRPPVLRAGVNTVTI LVENKKAQLVVIAHDVDPIELV VFLPALICRKMGVPYCIIKGKAS LGRIVHRKTCTTVAFTQVNSED KGWLAKLVEAIRTNYNDRYD EIRRHWGGNVLGPFKSVARIAK LEENAKAKELATKLGLNVHC
15277	45645	A	15367	2	1006	WINLIGNILEKRESLTSNFWPCK LASRKRHTTRNLEMNLWLSSH PQQSNITEWCCQLSCSTSLLT LLGTISLPLLSYLJHLIPPGPRQVY HRTSTLLLGHISVGEKFHKS QHWGFCNNVMMLQLQLSPLK GLSLVDKVNRPPALSGTRVLAS YEDEPLLRGNPCRFVIFIEWHD IWOMYEKAEASFWTTEEVDLS KDIRHWESLKPERFYEISHVLVF FAASDGINENLVERFSQIVQITE ARCFYGFQJAMENIHSEMYSLLI DTYIKDPKEREFLYNJETMPC VKKKADWALRWIGDKEATYG
15278	45646	A	15368	3	1368	QAQPMGRVGGMAQPMGRAGA PKPMGRAGSARRGFKGGCWSE GSPVHPVPAVLSWLLALLRCAS TMI.SLRVPLAPITDPQQLQLSPL KGGLSLVDREWTPPALSGTRVL ASKTARRIFOEPTEPKTKAAAP OFVEDEPLIRENPREFVIFIPIEYH DIWQMYKKAEASFWTAEEV\D LSKDIQHWESLKPEERYFISHVL AFFAASDGIVNENL VERFSQEV QITEAJRCFYGFQIAMENIHSEM YSLLIDTYIKDPKERFEFLENAIE TMPCVKKKADWALRWIGDKE ATYGERVVAFAAVEGIFSGSF ASIFWLKKRGLMFGLTFSNELIS RDEGULICDFACLMFKHLVHKP SEERVVEIIINAVRIEQEFLTEAL PKKLIGMKGTLTMKQYIEFVADR LMI.ELGFSKVFRVENPEPFMEN ISLEGKTNFPERKVGVGVMGV MSSPTENSFILDADF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15279	45647	Α	15369	2	621	RFIKMLSRHHQHVHHVLPLLRA
						PLPVAMEEEIPALFIDNGSGM/C
				i		KSSFA/GDNALRAIFPSIIGHPRH
						QGVMVGMGQKDSYVGDQAQS
				1		KCGILTLKYPIKHGIVTNWDDM
						EKIWHHVFYNELCVALEEQVV
				ļ		LLTEAPLNPRANREKMTQIMFK
						TFNTQAMYVAIQAVLTLHSSGC
						TTGIVMDSGDGVTHTVPIYERH
			1			TLPHTILHLDL
15280	45648	В	15370	1	966	
15281	45649	Α	15371	2	228	
15282	45650	В	15372	1	816	
15283	45651	Α	15373	256	384	RKLLNEAPC*PMR*HSLLNRLA
				<u> </u>		TSCNSWTVPVVPKFSKLRRGL
15284	45652	Α	15374	145	375	KIAGKNRLNRNQCKERKVQVL
				1		WISEKKKQREAVEDSPSLDCCH
						KFPPGS*ERLHG*KRGQWF*PP
		<u> </u>				HTVEKQMRSCLL
15285	45653	Α	15375	319	482	
15286	45654	Α	15376	4776	5910	VCSTAGDVNGVCLLYDLLHIA
	1	i				VASGSAKGECFFARKPLLAEGV
1	1			1		IYDCLQEFKEKKLVPATPHAQV
	i	ļ		l		LSYEALLSAHDTIAQKDFEPLLP
ŀ	1	1		ı		PLPDNIPESEEAMRIVCLVKNQ
l		1		1		QPLGATIKRHEMTGDILVARIIH
				ł		GGLAERSGLLYAGDKLVEVNG
1		1		i		VSVEGLDPEQVIHILAPRPCLSP
		1		l		MSLADLPPLPREVLVLAGKLPV
1				1		PSSSFLLQPREDCQGPQPSSLIT
		i		l		WFAPLETEPGVDRAMSRGTIMF
				1		KVVPVSDPPVNSQQMVYVRA
1				1		MTEYWPQEDPDIPCMDAGLPF
	į	l		1		QKGDILQIVDQNDALWWQARK
				1		ISDPATCAGLVPSNHLLK/RIWT
	l					EPGICIFTKRLRCSGKQREFWW
16007	45655	A	15377	2	405	SQPYQPHTCLKSTLCEYCNCPT WSLVTRRALARVGLPG/SPPPRL
15287	43633	l ^A	15377	ľ	403	LLLPLLLGWGLRVAAAASASSS
						GAAAEDSSAMEELATEKEAEES
						HRODSVSLLTFILLLTLTILTIWL
				İ		FKHRRVRFLHETGLAMIYGLEK
ĺ						
15288	45656	A	15378	1 -	3804	ITIGVLDITGWPYCTIQPYNGFL
15289	45657	A	15379	349	622	RPDVANLAFLARKEGTKRGFLS
.5207	1.5057	Ι΄.		["	I	KKTAEASRWHEKWFALYONV
				1		LFYFEGEQSCRP\GMYLLEGCSS
				I		GGTSPRSFRSSSRDSPISIPLHPT
						AK
15290	45658	Α	15380	314	483	
-5255	1.5000	1				

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15291	45659	A	15381	352	822	IDPRRVQAFPAKMKFNPFVTSDR SKNRKRHFNAP\SHIRRKIMSSP LSKELRQKYNVRSMPIRKDDEV QVVRGHYKGQQIGKVVQVYR KKYVIYIERVQBEKANGTTVHV G\IHPSRVVITRLKLDK\DRKRIS DRKAKSRQVGKEKGKYKEETI EKMQE
15292	45660	В	15382	80	432	
15293	45661	A	15383		888	VKCRKAEGRRISKLQTFEESQA VEAAMANVPWAEVCEKFQAA LALSRVELHKNPEKEPYKSKYS ARANMIEEVKALLGPAPEDED E ERPEAEDEPGAGDHALGLPAE VVEPEGPVAQANRLRLLAVIEF HLGVVHIIDTEELSAGEEHLVKC RLILRRYRLSHDGISLCIQAQT WRCIGVRDPPTSSCAFLSTSPTP VHSVKKIQTPGPLRPSNSTLVD DMESSPGTCASPLSQDVVNGS NQEEIGTGVDRCHKFLDIARQ NISFLQKRLQLSVQKPEQVINED VDVSELRNE
15294	45662	A	15384	1864	2705	WPGKVLAHRGLIPWIRDPHGS GGGAGQELRRQGCLEQLLYD AAHAQRLPPLDSVLPRAALPQD CQRASACEGDPTLLRPSHSPGH PRGAEER/PSTACLQRSWEGK WNRALQVGGLKSPWRGEYK EPRHPPPNQANYHQTL/ACPAE R/CFRQGPQGFQGLRRQQAEPL SSSLLSHQSPQSOTSLLP*L*AR RSLGGGNPYLCPPWSQPLFETP AHQSGKQPSRSRNCSSWK*NYS STACPSHFLWKRSKFSRASAST
15295	45663	A	15385	1113	1378	RPHAESFVSGGIPDPPGPGPGST GPGPL/ERQPCSLTSWPREPPSSS SMKMLTQGPSLTAPYKGTILGE VNPAHAISSSARNTSSRTSV
15296	45664	А	15386	285	722 .	TPAHPGVPFGQSGRHRHW/QSA ARSQLQPSAW*PKTGSLFATTW RCQTRASTCSAHWPLMAASPG AGGSSMASWR/DRP*PCPPPAP HLPEAAFLLR/CHDACP*NTGSA VPRGSCQPPAQQWDQGLAATR WGQAERLPRIYNWALAPP

SEQID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540.217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15297	45665	Α	15387	207	605	FQASAERFGTEDD/GSLRTRPPV
						MQSHQWPWRRPRPLVSRGRPQ
		l				TTPERHDSGGSLPLTPRMESHS
		1		1	1	EDEDLA/EGCRWP/EAGTVGVP
						GPRAQGAAQR/QAVLARKKHR
	ľ			i		RRPSKRKRHWRPYLELSWAEK
						QQRDERQS
15298	45666	Α	15388	193	407	ACPKTSKNLRQRRLWNLAKHE
						F/SVDMTCGGCAEAVSRVLISLE
l						VRVKYDIDLPNK/KVCIESEHS
						MDTLLATL
15299	45667	В	15389	321	625	
15300	45668	Α	15390	275	690	SLWPPVIRG/SPKGNRPAILTYH
	1	l				DVGLNMANKKSGDHPEREGTP
	1					RSFREVVSAFGPGSSKLQIPKGS
	İ	l				APRCGVPGRGARRELQSFRAGP
	l	l				RTQGSPVGPRCQSRRLRSAQEE
	İ	ŀ				SAKARHGGLRAHRSFSWDIRM
	15550	ļ.	1.5201	2	137	SSKERKFS PGRYRPN*LLLSDASPGAGNEF
15301	45669	Α	15391	2	137	
15302	45670	A	15392	1	648	QSTFLTPTILPSLRFPVLYTPNS MSLRKLTIMVEGKGGPNIPHAC
15302	43670	l ^A	13392	1	040	GCRORSSIKVSLLPAVTSKSESP
						KEPEOLRKLFIGGLSFETTDESL
	ŀ	1				RSHFEORRTLTDCAVMRDPNT
						KCSKGFGFVTYATMEEKYHTV
		1				NGHSCEARKALSKQEVASASSS
		1				OR/GRSGSGNFGGGHGGGFGG
	i	1				NDNFGHGENFRGHSSFGGSHG
1		l				GSGYGGSGDDYSGFGNDGSNF
		l				GGGGSYNDFGNYSNOSSNFGP
15303	45671	A	15393	1	3629	MTGICYTEDERSYKKNAOPTA
						ASKKQKETQKFCLRVDGQQKV
1						KLSVLOEKSAOLTVOLKSOKFL
l		l				GHPTAGRGRSELCLDLPDPPED
		l				PVALETRSVGTWVRERDLGMP
		l				DGEAALAAKVAVLETQLKKAL
İ						QELQAAQARQADPQPQAWPPP
1		l				DSPVRVDTVRVVEGPREVEVV
1						ASTAAGAPAQRAQSLEPYGTG
l		1		1		LRALAMPGRPESPPVFRSQEVV
1					1	ETMCPVPAAATSNVHMV KKISI
		1				TERSCDGAAEMKWEDQNIGD

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		E
		_				<u> </u>
15304	45672	Α	15394	1	1515	MGQVFRVVTGVGEQPGHPDQF
						PYIDSWLSIIQNHPKWLQACFET
		1		1		YCKTLMAQIKPGTIERDCKASV
		1		1		KEKDSQEKQKKPVLQAPPEELE
						SPALYAPIYPSLARLRQEAAPA
						APRGSDSEESTPQATPCREEPEP
ŀ						LPEKETSSSYRSNAVHFLIHNPT
			1			WADCKQLLQSLFNTEEHHRVI
	l					QAAFQWLENNAPAGTGYIRQY
						AQQALPIEADTGWDPNQAQGL
						QSLQWYREALLNGIKAGGKKA
						TNIGKVSEVCQKPDESPSEFYER
1						LCEAYQLYTPFDPEAAGNQCM
1		1				VNVAFVSQAQGDIKRKLQKLE
ł						AKAVLRGKFIALNTYIKKAERA
						QTDILRSHLKELKKQEQTKSKA
	ļ	1				SRRKDVTKIRAELNEIETTTTKI
İ		1				QKINETKSWFFEKINTINRPLAR
		1				LTKKSREKLQITSLRKEMGDTT
						TOTTEIQKIIQGYYEHLYAHKLE
						NLEEMNKFLEKYNAPSLNQEEL
1						HTLNRPIRNSEIEVVIKKLPTKK/
ŀ						SPGQDGFRAESYQIFKEALV
15305	45673	В	15395	1	1077	
15306	45674	Α	15396	69	290	YYFDSPIHPLHLSGTLRPGGSRQ
						FPCDPQFLCRLLHTLASL*SLPG
					1	DLLRGLLHKLVCRVERLDLASL
					1	YAVGR
15307	45675	A	15397	62	201	
15308	45676	Α	15398	3	429	LPSSVPRQPKMQARCLSRSIRA
	1			1		VEIYESDLEEGQGLLHEFSQFVL
i						DRPPPTQSRYAQELVHTCYLHT
						LASL*SLPGDLQRGLLHKLVCR
					i	AERLDLASHHAAVGRSELFFW
ł						TPYTHSPPGAGVGHPFPWHTRL
				İ		PWGVPAEGHQ
15309	45677	A	15399	303	612	PSSSIPVYHCSLRGICIPVLGLKR
	1	1				AANVGVSLEASPPLLSTPGGLG
1		1		1		CLSCGGTRTVGSRVGSIFPW*G
1	1	1		1	I	EGTTGITGSISCKGSSDVGSNRT
1		1		1	1	SGVDFPRRVE
15310	45678	A	15400	I	273	
	1.50.0	1	1.5.00	1	E	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15311	45679	Α	15401	1	406	MEEGWIRLPDGRVAAPQLLGA
						AVVLAVQETTHRGQESLEKLL
						GRYFYISPLSALAKTVRQRDFA
						DFGTTIKQDFRLLGQTSVDRLL
						QLSQGQAVKGNQLLPVSLTSQ
						CQVYKCVWNWWVLGLTDFKN
						EATDPPGVKLQTFLVSVTTHKG
						SVDPNSEKQQDLLQRAKEQSFH
		1				SVEGDPSRDFADFGTTIKQDFR
		1				LLGQTSVDRLLQLSQGQAVKG
		1	İ			NQLLPVSLVKRKTTLAPNTQTA
						SPRALADSLMQLARQVSR*RAI
						SCCPSH*LPSAKCISVSGIGGFL
		<u></u>				VSLTSRMKPRTLLE
15312	45680	В	15402	1	930	
15313	45681	A	15403	391	617	LQLQPTG/CNFLYFYVGYMKIL
		1				GFCFHHIVEERGGCLWYPCFYC
		i				NLSLLGG*RSCSHTRIVPSLLMP
		_				KLGPGGQVC
15314	45682	A	15404	1	1860	
15315	45683	A	15405	1	531	
15316	45684	A	15406	2302	2691	
15317	45685	A	15407	520 3690	2883 4727	ENGUING CEUDODIC COUL CDI A
15318	45686	A	13408	3690	4/2/	ENCHWGCEHPCDIGSSIILSPLA YWEOYHRVTYISRDTEGSIIVFP
						ALVTMLRNLFIILAGSSDPHFHT
		1				PMYFFLSNLSWADIGFTSATVP
		1		i		KMIVDMOSHSRVISYAGCLTO
						MSFFVLFACIEDMLLTLMAYDR
						FVAICRPLHYPVIVNPHLCVFFV
						LVSFFLSLLDSOLHSWIVLOFTF
						FKNVEISNFVCDPSOLLNLACS
						DSVINSIFIYLDSIMFGFLPISGIL
						LSYANNVPSILRISSSDRKSKAF
			l			STCGSHLAVVCLFYGTGIGVYL
l		1		I		TSA\VSPPPRNGVMASVMYAV
						VTPMLNLFIYSLRNRDIOSALRR
l						LLSRTVESHDLLSQDLLHPFSC
			ĺ	1		VGEKGOPH
15319	45687	В	15409	1	3195	
15320	45688	В	15410	1	1819	
	1.5000		1	1.	1	i e e e e e e e e e e e e e e e e e e e

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of tast	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15321	45689	A	15411	399	1089	LLTPRTDLQEMPLDNAEIEWYT
						DGSYLRGEVFQWYDHLALIFVP
	ŀ					QTGIEDVIWHIEALPNYIQKVCE
			ŀ			RRNPYNQILSLQCAQHEIFLVS
1	ŀ					WKLVCKQLEGDDDMLVGTTA
1				l		SNLLLESEQSTSNLNEKINHLEY
1		1		ŀ		EDQYKDNNFGEGNDGGILEDK
	ľ					LISN/NMGGPDSPDSVNPVE/P/M
						PTMTDQTTLVPNEEEAFALEPI
	İ	ı				DITVKETNAKRKRKLIVDSVKN
1						LDSKTIRAQLNDY
15322	45690	A	15412	3	809	FKDDDMLVSTTTSNLLLESEQS
		1		l		TSNLNEKINHLEYEDQYKVDNF
		1	İ	ł		GEGNDGGILDDKLISNNDGGIF
		l				DDPPALSEAGVMLPEQPAHDD
}		1	1			MDEDDNVSMGGPDSPDSVDPV
		1		l		EPMPTMTDQTTLVPNEEEAFAL
		1				EPIDITVKETKAKRKRKLIVDSV
		1				KELDSKTIRAHLSDYSDIGTTLD
				i		LAPPTKKLMMWKETGGVQKLF
			ľ	l		SLPAOPLWNNRLLKLFTRCLTP
						LVPQDLIKRRKGGEADNLDEFL
		l		l		KEFEHPEVPREDQQQ*HQQRD
				l		VIDEPII
15323	4569 I	Α	15413	1	595	MVNRSHFRGVGEPCHQQNSHG
		ł				HLNSITETSNSPTLRAAPLQPTR
		l				DPSDMAVFEIDELPEGAVKPPA
						NKYPIFFFGTHETAFLGPKDLFP
		l		l		YKEYKDKFGK\SNKRKGFNEGL
				l		WEIENN\PGVKFTGYQAIQQQS
		1		l		SSETEGEGGNTADASSEEEGDR
		l		l	1	VEEDGKGKRKNEKAGSKRKKS
				İ		YTSKVTKKLLIVINVICISIKHF
15324	45692	Α	15414	41	785	GKGWAGGASGPANHGPAAGG
		1				RSPGMLRGDQGLPGRGAAGM
	i		İ			ARPRSRE\YKAGDLVF\AKMKG
		l				YPHWPARIDELPEASVKPPTNK\
						YPIFFFWHPMKPAFLGPQ\DLFP
		Ì			İ	Y*EYRGKFGK\SNNRKGFNEGL
	1	l	1			W/EIENNPGS/IRFTGYQA/IQQQ
	1	ĺ		İ		SSFRTEGEGGNTADASSEEEGD
		1		1		RVERDGKG\KRKNEKAGSKRK
		1		l		KSYTSKKSSKQS\RKSPGDED\D
		1		l		KDCKEEENKSSSEGGDAG\NDT
		1		l		
	L		1	l		RNTTSDLQKTSEGT

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15325	45693	A	15415	1	1272	MWSGAGPGLAVWPQLVTVLSL
						SLSRLNCKWGLSFLACLLLGD
						MADAAMQHYGVNGYSLHAM
			İ			NSLSAMYNLHQQAAQQAQHA
						PDYRPSVHALTLAERLADIILEA
			İ			RYGSOHRKORRSRTAFTAOOL
						EALEKTFOKTHYPDVVMRERL
						AMCTNLPEARVQVWFKNRRA
			1			KFRKKORSLOKEQLOKOKEAE
						GSHGEGKAEAPTPDTOLDTEOP
						PRLPGSDPPAELHLSLSEGSASE
						SAPEDOPDREEDPRAGAEDPKA
						EKSPGADSKGLGCKRGSPKADS
						PGSLTITPVAPGGGLLGPSHSYS
						SSPL/SPLPSAGAIPPAHGGHQQ
						PGALLVLRSRGSGPCCCSGGCC
						CALPGROHGPAGLTA/SASPTTS
						PCOOPLLP/SOGVWGSPLLPAPP
						AGLAPASATLNSKTTSIENLRLR
						AKQHAASLGLDTLPN
15326	45694	Α	15416	1	1152	
15327	45695	Α	15417	8	82	
15328	45696	A	15418	1	1350	MGSEKDSESPRSTSLHAAAPDP
						KCRSGGRRRRLTLHSVFSASAR
		l				GRRARAKPQAEPPPPAAQPPPA
		1				PAPAAAQGPPPEALPAEPAAEA
		1				EAEAAAAAAEPGFDDEEAAEG
		1				GGPGAEEVECPLCLVRLPPERA
		1				PRLLSCPHRSCRDCLRHYLRLEI
				l		SESRVPISCPECSERLNPHDIRLL
		1				LADPPLMHKYEEFMLRRYLAS
				l		DPDCRWCPAPDCGYAVIAYGC
		i				ASCPKLTCEREGCQTEFCYHCK
						QIWHPNQTCDMARQQRAQTLR
						VRTKHTSGLSYGQESGPADDIK
		l		l		PCPRCSAYIIKMNDGSCNHMTC
						AVCGCEFCWLCMKEISDLHYL
	1	1	1			SPSGCTFWGKKPWSRKKKILW
		1	1			QLGTLIGAPVGISLIAGIAIPAM
	1	1	1			VIGI\PVYVGRNIHSRYDGKEN\S
	1	1	1			KHKRNLAITGGVTLSVIASPVIA
		1	1			AVSVG\IGVP\IMLAYVYGVVPI
		1				SLCRGGGCGS
15329	45697	Α	15419	2	540	

	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
15330	NO:	of peptide	hod				
15330		sequence	1	09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
GGGGPTRKPPRRTSSHLPSSPA					sequence		
GGGGPTRKPPRRTSSHLPSSPA	15330	45698	A	15420	1	530	SRTTPGK APKETWGODAGSGK
AGDCPILATPESPPPLAPETPDE	15550	1.5070	ľ.,	15.00	ľ		
AASVAADSDVQVPGPAASPKP			l				
LGRLRPRESKUTRRI.PGVARPE		ı	1				
AGDGHLSAVAERPKVSLHFD		ŀ	1		ĺ		
TETDGYPSDGEMSDSDVFAED			1		i		
GGVQRGPREAGAKE\VVRMGV		i	1				
LAS			1				
15331 45699 A 15421 1 2989 MILVMYWAPAPPPVHMSQFLT TGPKCLEISLDHGRDSGEITAS GILWQHAAGGLAQWNQKPH PPFLTLLQAEEPFERARSCDIVIF IKMQTAQGHQVAQFERRE ALRGSQSPQKSASQQRGSLDR ELLMLIKISKCLRVPSGPSSVLG TESTGPVPGELVLCQQDAGAT EGVCYGRELMICRCDSSPRLME PWGVKGTFEVFRDLITAMKJP DSYQLSPDDYYLLADPWRQEW EKGVQVPAGAEAIPE 15333 45701 A 15422 218 404 470 LTMLFAACVFSGDEIMASEERQ PMEEEWSRSSTISGEDGERWTP DTAGKAQGIAGDVWVSPLP PSKVSAASKPPKSIHHPFFKWH NFSNLMKMFYKMYTSKN TFENTRIFWDVDYPTGERRJE WALPPPGAGSVPAGGWSRPLTI GRCDSQGRGRSRAJE SSPL PPLRRSPTLPGEPPPAHRPLRVH PRHOPRPSHLVACLNSQAPDASR LRASANQ LRASANQ			1		i		
TGPKCLELSLDHGRDSQEITAS GLLWLQHAACGLAQWNQKPH PPETLLQAEEPFERASCDIVE IKMLQTAQQHQVGLAPTERKE ALRGSQSPQKSASGQRFGSLDR ELLMLIKISKCRVPSGPSSVLG TESTGPVPGELVLQQDAGATT EGVCYGRELMICRCDSFPLME PPGVKGTPEVFRTDLITAMKIP DSYQLSPDDYYILADPWRGEW EKGVQVPAGAEAIPE EKGVQVPAGAGSVBHPFFKWH NFSNLMKMFKWKYFSKN EKGVQVPAGGSVBHPFFKWH NFSNLMKMFKWKYFSKN EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLISSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGRSRAPLSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQVPAGGSVBPLTI GRCDSQGGRGSRAPALSSEL EKGVQTAT	16221	15500	 -	16421	,	2000	
GLLWLQHAAGGLAQWNQKPH PPFLTLLQAEEPFERARSCDIVIF IIKMLQTAQQHQVGLAPTERKE ALRGSQSPQKSASQQRFOSLDR ELLMILKISKCLEVPSGPSSVLG TESTGPVPGELVLQQDAGATC EGVCYGRELMICRCDSFPRLME PWGVKGTPEVPKTDLITAMKIP DSYQLSPDDYYILADPWRQEW EKGVQVPAGAEAIPE EKGVQVPAGASVPFKSIHPFFKWH NFSNLMKMFYKMYSKN TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGGRGSRRAAPLSSRPL EKGSQGGGRGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGRGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGRSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGRSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGGSRAAPLSSRPL EKGVQVPAGGWSRPLTI GRCDSQGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	13331	45699	I ^A	15421	['	2989	
PPFLTLLQAEDPFERARSCDIVIE		i	1				
IKMLQTAQQHQVGLAPTERKE ALRGSQSPQKSASQQRPGSLDR ELLMILIKISKCLRVPSGGSVLG TESTGPVPGELVLQQDAGATG TESTGPVPGELVLQQDAGATG EGVCYGREMICRGDSSPRLMB PWGVKGTPEVFRTDLITAMKIP DSYQLSPDDYYILADPWRQEW EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAEAIPE EKGVQVPAGAGAVFSKN EKGVQVPAGAGSVPAGGWSRPLTI GRCDSQGGRGRSRAPLISREJ EKGVQVPAGGSVPAGGWSRPLTI GRCDSQGGRGRSRAPLISREJ EKGVQVPAGGSVPAGGWSRPLTI GRCDSQGGRGRSRAPLSSREJ EPLRSPTLIGEPPAHRPLRVH PRHGPRPSHLJACLNSQAPDASR LRASSNQ LRASSNQ LRASSNQ EKASS		l					
ALRGSQSPQKSASQQRGSLDR		1	1				
ELLMLİKISKCLRVPSGPSSVLG TESTGPVPGELVLCQQDGAGTC EGVCYGRELMICRCDSFPLLME PWGVKGTFEVFRIDLITAMKIP DSYQLSPDDYYILADPWRQEW EKGVQVPAGAEAIPE 15333 45700 A 15422 218 404 15333 45701 A 15423 1 422 15334 45702 A 15424 160 470 LTMLFAACVFSGDEIMASEERG PMEEWSRSTISGEDGERWTP DTAGKAAQGIAGDVVWVSPLP PSKVSAASKPPKSJHFPFFKWH NFSNLMKMFVKWYFSKN 15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGRGRSRAAPLSRSPL PPLRSPTLPGEPPAHRPLRVH PRHGPRPSHLVACLNSQAPDASR LRASANQ LRASANQ							
TESTGPYPGELVLQQDAGATC EGVCYGRELMICRCDSFPRLME PWGVKGTPEVPKTDLITAMKIP DSYQLSPDDYYLLADPWRQEW EKGVQVPAGAEAIPE 15332		l					
EGVÇYGRELMICRCDSFPRLME PWGVKGTPEVFRTDLITAMKIP DSYQLSPDDYYLLADPWRQEW EKGVQVPAGAEAIPE 15332 45700			1		Í		
PWGVKGTPEVFRTDLITAMKIPDSYQLSPDDYYILADPWRQEW EKGVQVPAGAEAIPE 15332 45700			l				
DSYQLSPDDYYILADPWRQEW	İ						
EKGVQVPAGAEAIPE			l				
15332 45700 A 15422 218 404 15334 45701 A 15423 1 422 15334 45702 A 15424 160 470 LTMLFAACVFSGDEIMASEERQ PMEEEWSRSSTISGEDGERWTP DTAGKAAQGIAGDVWVSPJP. PSKVSAASKPPKSIHFPFFKWH NFSNLMKMFYKMYFSKN 15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGGRGRSRAPLSRSL PLRASHAPLSRSL PLRASHAPLSRSL PLRASHAPLSRSL PLRASHAPLSRSL LRASSANQ LRASSANQ			1				
15333							EKGVQVPAGAEAIPE
15334 45702 A 15424 160 470 LTMLFAACVFSGDEIMASEERQ PMEEEWSRSSTISGEDGERWTP DTAGKAAQGIAGDVVWSPLP PSKVSAASKPPKSHIPFFKWH NFSNLMKMFKWMFSKN 15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGGRGRSRAPLSRSPL PPLRRSPTLPGEPPAHRPLRVH PRHGPRPSHLVACLNSQAPDASR LRASANQ LRASANQ LRASANQ							
PMEEEWSRSSTISGEDGERWTP DTAGKAAQGIAGDVVWVSPLP PSKVSAASKPRESWHPFFKWH NFSNLMKMFYKMYFSKN 15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGRGRSRRAAPLSRSPL PPLRSPTLPGEPPAHRPLRVH PRHGPRPSHLACLNSQAPDASR LRASANQ LRASANQ					·		
DTAGKAAQGIAGDYVWVSPLP PSKVSAASKPPKSHHPFFKWH NFSNLMKMFVKMYFSKN 15335 45703 A 15425 21 375 TFENTREWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGRGSRRAAPLSRSPL PPLRRSPTLPGEPPAHRPLRVH PRHGPRPSHLACLNSQAPDASR LRASANQ LRASANQ	15334	45702	Α	15424	160	470	
PSKVSAASKPPKSUHPIPFKWH NFSNLMKMFYKMYFSKN 15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDQGGRGRSRAAPLSRSPL PPLRRSPTLPGEPPAHRPLRVH PRHOPRPSHLVACLNSQAPDASR LRASANQ LRASANQ			!		ì		
15335 45703 A 15425 21 375 TFENTERWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSPRJTI GRCDSQGRGRSRRAAPLSRSPL PPLRRSPTLPGEPPPAHRPLRVH PRHGPRPSHLACLNSQAPDASF LRASANQ		1	1				
15335 45703 A 15425 21 375 TFENTRFWDVQIPVPTGERRNE WALPPPGAGSVPAGGWSRPLTI GRCDSQGGRGSRAAPLSRSLL PPLRRSPTLPGEPPAHRPLRVH PRHGPRPSHLVACLNSQAPDASR LRASANQ		1					
WALPPPGAGSVPAGGWSRPLTI GRCDSQGRGRSRAAPLSRSPL PPLRRSPTLPGEPPPAHRPLRVH PRHGPRPSHL\ACLINSQAPDASR LRASANQ							
GRCDSQGRGRSRRAAPLSRSPL PPLRRSPTLPGEPPAHRPLRVH PRHGPRPSHL\ACLNSQAPDASR LRASANQ	15335	45703	Α	15425	21	375	TFENTRFWDVQIPVPTGERRNE
PPLRRSPTLPGEPPPAHRPLRVH PRHGPRPSHL/ACLNSQAPDASR LRASANQ			1				WALPPPGAGSVPAGGWSRPLTP
PRHGPRPSHL\ACLNSQAPDASR LRASANQ	l	1		1			GRCDSQGRGRSRRAAPLSRSPL
LRASANQ	1	1			l	1	PPLRRSPTLPGEPPPAHRPLRVH
			1	l	1	1	PRHGPRPSHL\ACLNSQAPDASR
15336 45704 A 15426 155 340	l			1	1		LRASANQ
	15336	45704	Α	15426	155	340	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15337	45705	A	15427	sequence	2658	MIINTITTTTTITIFIITIMSSPIT ITNIITIOHHHHHHDQRPSLGK CRGLYIRVRKVPGERATANSRC PINAQGIESCWNGVMALWLGP STCSRLPELTDSEDCQPADSP GDEDPKPWWDDPHEAPLTGYF HPYGAAFVPRHAGNFKDRRBV PHQDTSFCSAGGFWILPSACKT GRVALTLTLLIGAYQTLTEADV KGLTAARAASPGAKWRTQFFC PAGTCSSQAGNGQLEDCLPYPP GPSAPVGSLSRSANGEFIIGRVI KAMNNSWIPECFRCDLCQEVL ADIGFVKNAGRHLCRPCHNSE KARGLGKYLQCKHAIIDEQFEL KRAGLGKYLQCKHAIIDEQFEL FKNDPYHPDHFNCANCGIKLG VFLTDRKDLTADAQELKGELY VFLTDRKDLTADAQELKGELY CLPCHDKMGVPIGGACRRPIEG RVVNAMGKQWHYEHFVCAKC EKPFLGHRHYERKGLAYCETH YNQLFGDVCFHCNRVIEGDDE
						GOĞFAHRYSPYDSCEVDQSPLE GQQLKSIRKGPLAQSQPLPTPY MLHPGCKTQIREAPGSAALESV RRPRPRGL VPEESSESAAVEAV RRHHLQNFSSLJETLINGQSLG GHFCDVTVCHHEASILATINGSLG GHFCDVTVCHHEASILATINGSLV VPVQTVRQLVELLYSGSLVVA QSEARQVLMAASVLRIDSYRR MHADYRTRSSAPASTSAPAPLP TPVPPPLAPEQLRHRLRHLLAA
15338	45706	Α	15428	708	892	
15339	45707 45708	A	15429 15430	212	713	DLPLAVHGQVSDSGAQAVASR SGDESSIVCLKEESTVMTFRS SGDESSIVCLKEESTVMTFRS GREVIKAMINSWHPECSAVTSA RKF WQLIHLSRMLVDRKDLTA DAQELKGELYCLPCHDKMGVP IGGACRRHEGREVVNAMGKQW HVEREQVMPDGILSALIDPKYL LILSHRGLHMKSMKGFYFAKL YYEAKEYDLAKKYICTYINVOG RDPKAHRFLGLLYELEENTEKA VECYRSVELNPTQKDLVLKIA LLICKNDVTDGRAKYWERA AKLFPGSPAIYKLKN/DGWVRG QMVPKPDDEGVQVRGRSGRPD AGWKLMRSSYGGSAWRLRPQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15341	45709	A	15431	129	444	WGGPGRQPDGGRVPGRCGPRR PATWESPAACRPRPQSRHHSFE
						*GGSRSGSSHKRPCPSSPAVLRA RDQVVLHTSFEATEGDSSHEDR
15342	45710	A	15432		2446	STPLARRCLCSHRSGEA METSLMWQVCGGRDCNALLPG NLCSQPTQVLPLEVDGDDGPG GLPLNAAPPSSVPVPPSEALLPP PACPSAPAPWRSIISRLFGTSPA AEAAHPPPGGALDFGALRTSSC LSQSHGHIPGGPGLGLGAVRVP VPSEEDPASLFTEPVPAAEAPAT VQSVEDFVANDRLDRSFLEDM TPARDEKK VGAKAAQODSDTC RASRAPEQQIPHGGHWRGHSRS YERFRPDMVSAGPGPSDGEAL GGNPIVAGFQDDVDLEDGPRG PPPPAGPVSQDITLSSEEAEV ADPPKGPAPAPQCSEPETKWS SPLASKEPGGASSESDPEGPHAAQML SFVMDDPPGESEASDTQCRAVF RVLPAQRAENAYGSAFQPGAL RVLPAQRAENAYGSAFQPGAL RVLPAQRAENAYGSAFQFGAL
						TEATSPRPTGGASPLAHGATVG AAWGLGMASWPIGSSLAPTRA RTREPRLQRRVGSELPGDPCSG SGASCRQVCARGRATSVRLPAL
						ESLWPLSSDGSAAAREGLEGPR LRQAAAPGGPGPPLPPLSRPPSP ARRMRGGAAVAGRGAAAPLE GRVRLGVGFLAETAVRAGAGA LAAADAPPGSRTPRARPLVAAC
						SGAAGGRCPVPSRRQREAPSPL GGAGVDLQLECLTSEKAGVGA ATGGVLEALARAIRQKKEIEEIQ

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SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15343	45711	Α	15433	26	963	FMIWAGQNRKVQDFVIQLRML
						TPVVPPPQAAPNAQQPRAAASA
						PQTRVRAPRWPLGNVVVLGTE
		ļ				DAAGPSRPASEPPASPALPGPAL
		1				ELGLPNHPKQT/HRPRLAPRRSH
		1				PKDSCTIFKLNSFLKNIKFNNSV
		1		i		GDQVNLDYKLISGTEYDILNFW
		1				NFRILAKTGCSKDGHRNPRSRP
		l				TQEEGSGKPKKSLVKERSGAQE
		i				DLPVPPEEKLELSGTSMGPCCFS
		1				KTSYGIPHIIYILSYKGTTATAG
		1				HRWPAAYPAPLNCSVPPPETTM
		1				PVSDLGSPLHWASRIATPPSMW
						LCCLLAAALRVLAVSNSSAEVY
		1				sqs
15344	45712	Α	15434	803	1305	RSKPCRSSHNVFSWWAHSRPC
		l				CPWSGYSSRLCCLSCCITLGPY
		l				QLKNRQQRNGSYKSFGNYLCV
1		l				GNRLLFWVWIIFQGKGRS/WLP
						RSDWVTSYKVMVSNDSHTWV
						TVKNGSGDMIFEGNSEKEIPVL
						NELPVPMVARYIRINPQSWFDN
	l					GSICMRMEILGCPLPGWEH
15345	45713	A	15435	2	377	ARELLKAVWRPHKVAVMHCR
	1					GHQRASTLVGLGNSRADSEAR
	1					KAPSAPFRASVTAPLLPQAPDL
						VPTYSKEENDFLQAKGGQVME
						EG*IRLPDGRVAAVVLAVHETT
		_				HRGQELLEKLLGWYFYISH
15346	45714	A	15436	77	679	KMKTNQMGLKSKVVEMNATIS
						EIRLNINRKDVEYQQEILQLEAV
		1				WKLHKVAVMHCKGHQRASTL
ŀ		1				VGLGNSRTDSESPKAASAPLRA
		1		1		SVTAPLLPQAPDLVPTYSKEEK
1			1	1		DFLQVEGGQVMEEGCIRLLDG
		1				RVAVPQLLGAAIVLAVHETTHL
		1				RS/ESSLGKLLGQLFSTISHFVSL
						CQNLTQRCVTCRQHNARQGPA
	l	<u>L_</u>				VPPGMQ

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for tast amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15347	45715	A	15437	2	881	HARCTGIAARAGLLDPRHTGEA
13317	13713		13.57	ľ	001	LAARSRIDLARRPHWPRGCGTC
						GGGGGCAGLGGAWLRGNDVA
	1	ļ				VPARALVTAAPASSSMSPPIPA
	1					AARPVSSS\PAASPSPRCGLRAG
ŀ						
						PAELRCPRRLRQARAEEPPRAG PAPGRRRPTPGICPAPLSPRLRA
						GGTAPSLPGVTRGSRVGRRLRP
	i					SLGGAGCRVGKRPPRLLPPGRR
						CSTGLPPPAGTGAPAGVCWED
	l					GDPGHELAAALLRRLGGQIRAL
						AGHRTAGWAQHGLSPREAVRP
						APAARGTLLLRRSNSRRHLQNP
		l				ANFGGTLAT
15348	45716	A	15438	3	502	GRRDSCLPGVVAPRACLHPRG
						QVPRLGSVGKMATPGMSWQQ
1		1				HYYGGSAAKFAPSPATAQLAG
						HSMDYSQEMHLKMSKKIAQLT
1	1	1				KVIYALNTKNDEHESAIQALKD
1	l .	1				AHEEEIQHILAETREKI\LQYKS
	l					KVTEELDLRRKIQVLESSLEDHI
		┞				KMKQQALTEFEAYKHR
15349	45717	Α	15439	29	230	LQIDCKNKPAILRGPTDPLKEA
						DCYCRTWERPQILLPKWEGTR
		L				KPILVI*QNKAHQHAPQITLLHQ
15350	45718	Α	15440	73	335	FFWGGIEEPCFVILSGSVFWFLL
	l	1				IWVGASASPVVGDVFGRGGSPF
1		1			1	PTSTVGAFTVLWGVSWVLKRV
15351	45719	A	15441	373	513	CGSSWDC*FVLAINLELKFTM
15351	45720	A	15441	1585	1649	KNSNIOHSPKITLVHOOWFOTK
15552	43720	^	13442	1303	1049	KKFLIYLKKNSGGLVDP*MVHT
l	l			l		TRLCADNPQYQPRARKSHS*VK
				i		GESTTLREHPMGQKNLNNSLQP
15353	4572I	В	15443	1	696	GEST TEREHFMOORNENNSEQF
15354	45722	В	15444	49	525	
15355	45723	A	15445	113	474	SIPYKNKEEEEGKGGEYYIKGT
13333	43723	^	13443	1113	474	CHGTKGSEOOPSALDLPSDRAY
				i		PNEKEPEN*LW*LI*QNKAL*HS
		i	ĺ			PKNHTSSP/RNGSKPRKKSLINL
						l
						KKNSGG*LLS*LGRHQRKVKPN
15356	45704	 	15446	47	674	ARKSKNQYKK
15356	45724	Α	15446	47	614	TGCPRCKQNSTCIAAVKMEGPL
				1		SVFGDRSTGETIRSQN\DVTITN
	l			1		DGATILKLLEVEHPAAKVLCEL
1	1	1	l		1	ADLQDKEVGDGTTSVVIIAAEL
				I	l	LKNADELVKQKIHPTSVISGYR
		1	1			LACKEAVFYLTENLIVNTDELG
	1					RDCLINAAKTSMSSDIIGINGDF
	1	1				FAN/MVVDAVLAIHARHRGQ/P
				L	L	RYPVTLLY*SHGES

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequence		
15357	45725	В	15447	707	1826	
15358	45726	A	15448	108	397	
15359	45727	A	15449	1726	2464	SKAVQLEAGGLLQLFCCKMAP
13339	43727	<u> </u> ^	13449	1,20	2404	DPNITDVFGRTALHYAVYNEDT
				l		SMIEKLLSYGANIEECSEDEYPP
				l		LFLAVSQRKVKMVEFLLKKKA
	1					NINAVDYLGRSALIHAVTLGEK
l	1			l	0	DIVILLLQHNIDVFSRDVYGKL
				l		AEDYASEAKNRVIFELIYEYER
1				ŀ		KKHEELSINSNPVSSQKQPALK
1				l		ATSGKEDSISNIATEIKDGOKSG
						TVSSQKQPALKDTSDKNDSVSN
						TATEIKDEQKSGT\VSSQKQPAL
ļ				İ		KDTSDKNDSVSNTATEIKDEQK
						SGTSVFSETTGLEGYK
15360	45728	Ā	15450	2	675	
15361	45729	A	15451	ī	467	DHSSSPAREONWMEN\EFDELT
				1		EVGFRRWVITNSSELKKHVLTQ
l	ŀ]	ŀ		CKEAKNLEKRLGELLTRITSLE
1	ŀ		İ			KNINDLMELKNTARELHEAYTS
1						INSWINQAEERLSEIEDQLNEIK
1	l					REDKITEKMKNSTTVRVAASM
						QSKLLQGVAEEGPLRLTRSASF
15362	45730	Α	15452	2	632	
15363	45731	В	15453	355	894	
15364	45732	В	15454	99	716	
15365	45733	Α	15455	1	1140	
15366	45734	Α	15456	1	1476	MEVNREKQLNELEVIGSEEQNL
		l				EEEGLMIGGVAVRLVPDDIVIP
1		l				GGVNATNGTEARDALRVKVA
		1				MSVTLLSGIIQYLLSALGWSYY
						TVDGVSQKNPRALGVTADQLH
						AIFTTMSDEQASFRFGFVAIYLT
					1	EPLVRGFTTAAAVHVFTSMLK
l						YLFGVKTKRYSGIFSVVYSTVA
ł						VLQNVKNLNVCSLGVGLMVFG
	į .			i		LLLGGKEFNERFKEKLPAPIPLE
	1					FFADHNSSPAREQK WMENEFD
						EWTEVSFRRWVITNSSELKEHI
						LTQCKEAKNLEKRLEELLTRITS
l			ļ			LEKNINDLMELKNTAQEFHEA
	l					YTSINS/RNQTEERVSEIEDQLN
1						EI*CKDKIR\EKKRMKRNEQSLQ
	1	1		1		EIWDCVKRPNLRLTGVPESDGE
	ŀ	1		i		NGTKLENTLQDIIQENFHNLAR
1				i		QANSQIQEIQRTPQRYSSRRATP
	l	1		I		RHIIIRFTKVEMKEKMLRAARE
	l	1		l		KGRVTHKGNPIRLIADLSADTL
	45805	١.	15450	ļ	Line	QARRQWEPIFNILKENFQTRISH
15367	45735	A	15457	1	1185	
15368	45736	Α	15458	I	2367	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15369	45737	Α	15459	1	744	
15370	45738	В	15460	337	1515	
15371	45739	Α	15461	1	1197	
15372	45740	A	15462	1	714	
15373	45741	A	15463	142	601	HQRTKVYKFTKMRKNQPKKPE NSK/TQNPSSPSKDINSLPAREE NWMENEFDKLTEVGFRRWVIT NSSELKEHVLTQCKEAKNPDK RLEKLLTRIISLEKNINDLVELK NTA*ELLEAYTNINSQIDQVEEK
15374	45742	A	15464	153	474	I/S/SIEDQLNEIKLEDEIREKKKK QTKSNNVNIKKQDVHTKTPSEG HQHQRPK\DHNSLPAREQNWIE NEFDELTEVGFRRWVITNSSEL KEHVLTQCKEAKNLEKRLQEL LTRITSLEKNINDLMELKNTA
15375	45743	Α	15465	1	2172	
15376	45744	A	15466		1640	IAKLMETL NOCINAGHENTKAI AIAQFNDDSPEARKITRAWRIG AIAQFNDDSPEARKITRAWRIG LPHPDMEIRGRVEQRVGYTIEQI GVAAHKGGVVKTSVSVHLAQ DLALKGLRVLLVEGNDPQGTA SWYHGWVPDLHIHAEDTLLPF YLGEKDDVTYAIKPTCWFGLDI IPSCLAL HRISTELMGKFDEGKL PTDPHI MLRLAIETVAHDYDVI VIDSAPNLGIGTINVVCAADVLI VIDSAPNLGIGTINVVCAADVLI VIDSAPNLGIGTINVVCAADVLI VIDSAPNLGIGTINVVCAADVLI VIDSAPNLGIGTINVVCAADVLI UVTPAELFDYTSALQFFDMLRD LLKNVDLKGFEPDHLTPAGNG GVYURREDGIGFFSGFYN VISL GVYLVEQOKGWEDTHSKTPSK GHOHQRSKVDKSTKMRCNWS KNAENSKNQNALSPRNGNSSP ARAQNWMENKFDKLTEVGFR ROYITNTSELKEHILTOSKEAKN LDKRLOELLTRITSLEKNINDL MELKNTARELREAFTSFCVNCE AVSLITSLREDSHVLEDSKGHR M/RRHQE*KCEYFHPPHFGLQF VQLPESDSINSDPLWPHGSGSLS
15377	45745	В	15467	50	1603	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, -possible nucleotide insertion)
				sequence		
15378	45746	A	15468	528 .	1413	QTERNSTSTSKTPSEGHOHORP
		l				KVDKSTKMGRNQCKKA*\NSK
		ı				SONACCPPKDHSSLPAREOTW
		ı				MENEFDELTEVGFRRWVITNSS
		l				KLKEHVLTOCKEAKNLEKRLE
		ı				ELLTRITNLEKNINDLMELKNT
		1				ARELREAYTSINSRIDQAKERM
		l				SEIVDOLNERNDRENGTKLENT
		l				LQDITQDNFPYLARQVNIQIQET
		1				QRMPQRYYSRRVTPRHIIVRFA
		1				KVEMKEKMLRAAREKDTSVK
		1	1			QKVNKDIQELNSALHQVDLIAI
		1				YRTLHPKSTEYTFFSAPHHTYS
		1				KTDHIVHSILCDSG
15379	45747	Α	15469	368	679	SSQLNKAPSLFNPVSEEFCLWL
		1				LLLQKELLTRITSLEKNTNDLV
		1		ļ		DLKNTA*ELHEAHTSINS*IDQV
		1				EERISECEDHLTEIRHSEKMKTA
						LLK*DMQTRREKKK
15380	45748	Α	15470	865	2055	QTERNSININKKDNYAKTPSKG
		l				HQQQRPKVDKSTKMRKNQHK
		l				KA*KFPKPGMPSSPPK\DHNCSP
		ļ				ARE\QNWMEN\EFNEFDELTGA
		1				GFRRWVITNSSELKEHVVTQCK
						EAKNLEKRL\QELLTRITSLEKN
						INDLMELKQDITQENFMKHTQ
	1					VSIAQADQGRKERV\SEIEDPTL
		l				MDIKCEDKIREKK/IKKRNEQSL
1	1					QEIWGYVKRPNLRLISVPESDG
						ENGNK*ENILQDIIQENF/PFPNL
		1				TRQANIQIQETQRTLQRYPLRR
	l					ATPRHIIRFTKVEMKEKMLRA
	l	1				AREKGRVTHKGKPIRLTENLSA ETL*ARRE**PILNILKERNFOPR
		l				ISYRAKLSFISK/GKREIKSFIDK
1		l				QTLRDFVTTRPALQELLKEAVN
		l		l		MERKNOYOPLOKHTKLKRPLT
15381	45749	A	15471	1	315	THE REAL PROPERTY OF THE PERSON OF THE PERSO
15382	45750	A	15472	i	2775	
15383	45751	A	15473	i	1017	
15384	45752	В	15474	i	3105	
10004	1.5.52		1.5 4	ı:	3.05	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15385	45753	A	15475	1	1115	MKWLLLIGLVALSECIMYKVP
13303	45755	<u>'`</u>	15475	ľ		LIRKKSLRRTLSERGLLKDFLK
						KHNLNPARKYFPOWEAPTLVD
						EOPLENYLDMEYFGTIGIGTPA
l						ODFTVVFDTGSSNLWVPSVYCS
						SLACTNHNRFNPEDSSTYOSTS
1				1		ETVSITYGTGSMTGILGYDTVO
						SGYHHGNHRLNLKGLCSSKRSS
						TTPPLYDLGQVASPPSGSVSSC
		1				DDOSGSVVIFGGIYSSYY\TGGL
						NWVPVTVEGYW\QITVDSITMN
	ľ					GEA\IACAEGCOA\IVEPGTFFVT
						GP\TSPIA\NIQSDIEA\SEELDG\D
		1				SVLICSAISS\LPDI\VFTI\NGVQY
						PVPPS\AYILQSEGSCISGFQG\M NLPTESEELWILGDVFHPASTLT
		ł				VF\DRANNOVG\LAPVA
15386	45754	A	15476	1	765	VFIDRANNQVGILAPVA
15387	45755	A	15477	470	479	PRNPSSFLQVQVQHRFLQLFVP
13367	43733	^	13477	470	4/7	FHFVFFLASLYVMVTLTTWFR*
		l				DPCVSGMALNCSFLKIASSSGE
	1	l l			i	YLRPTPSYLNOVVRVTMT
15388	45756	A	15478	1	896	CLPFGGPKPNPWDKDGGCOGR
13366	43736	^	13476	l'	070	POPRHLPGPGTAAQRRQQCPSE
		l			i	KSLL/CRWPSLTASTCSRLFYIL
		l		i		LHVGAS/CNLLPPAVKDSSGKG
1		l				VGQDTQVLSGSGAVYRVCAGT
1		l			İ	ATFHLLQAVLLVHLHSPTSPRA
		l				OLHNSFWLLKLLFLLGLCAIAF
		l				CIPDEHLFPAWHYIGICGGFAFI
1		l				LLOLVLITAFAHSWNKNCNEAS
1				I		YLAEVFGPLWIVKVYSYEFQKP
1		l	1	1		SLCFCCPETVEADKGQRGGAA
		l				RPADQETPPAPPVQVQHLFLQL
1		l		l		FCLPLRLLPWLTLCHGYPYQLV
I	I	ı	1	I	I	QASAGCGCKPFAPYSL

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15389	45757	A	15479	1	1016	MERRGPGAATARGRARPGGGP
13369	43/3/	^	13479	l'	1010	SVGLLATGSSLNPSFHGVARIVP
1					\	GFIRIARPRDGSFAYESVPWOOS
1						ATQPAGSLSVVTTVWGVGNAT
l	1					QSQVLGNPMGPAGSPSGSSMM
						PGVAGGSSALTSPQCLGQQAFA
						EGGANKGYVQQGVYSRGGYP
		1				GAPGFTTGYAGGPGGLGLPSH
						AARPSTDFTQAAAAAAVAAAA
		l				ATATATAT/GHRGCSPGEAEPG
						AEP\DGAMGAGQSFNSQFLQHG
		1				GPRGPSVPAGMNPTGIGGVMG
	ľ		l			PSGLSPLAMNPTRAAGMTPLY
				l		AGQRLPQHGYPGPPQAQPLPRQ
						GVKRTYSEVYPGQQYLQGAYP
		l	İ			VHSGLSQFPTWQETPRHP
15390	45758	С	15480	1	585	
15391	45759	В	15481	1	885	
15392	45760	Α	15482	3	440	GPWPRPVAYLSKQLYRVSKGW
1	1					PPGLRALAEMALLAQEADKLT
1	1					LRQNPNI/IGPHAVVTLMTTKG
	ł					HH*FTNARLTYVKIPT*PLKF/G
	1		l			NTLNPTTLLPVSESPVEHNCVD
	1		Į.			VLDSVYSSRPNLRDHP*TSVDC
	1	İ		l		ERYVDRSSFTNRCKVTR
15393	45761	Α	15483	3	368	
15394	45762	С	15484	109	276	
15395	45763	Α	15485	1	1710	
15396	45764	Α	15486	221	784	QNSLRSMCQKEKKMAAGVLT
						HTC\GPWPRPVAYLSKOLDRIS
						KGWPPGLRALAATALLAOEAD
						KLTL\GKT*I*RPPHAVVTLMNT
						KGHHWLTNARLTKYOSLLCEN
	İ		l			PHITTEVCNTLNPTTLLLVSESL
	1		İ			VEHNCVEVLDSVYSSRLNLRD
		ı				HP*TSVDW*LHVDGSSFTNPCK
		l				VTLKKMTSAAPVTPRS
15397	45765	В	15487	509	661	, I DICE, MI DICE VII KO
10097	CO 1 CT	<u> </u>	15707	1207	1001	

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
15398	45766	А	15488	1370	2243	LGKLRLESCGLVAAVRPHSGGL
		1		l		AMPLPNATRPASGRSED/AELK
1		l		l		LALEHGHRSESLQGRKHRGVH
						DNSDIPQAALVGGTTMIIGHVL
						PDKETSLVDAYEKCRGLADPK
ŀ		1				VCCDYALHVGITWWAPKVKA
		l		l		EMETLVREKGVNSFQMFMTYK
		ĺ				DLYMLRDSELYQVLHACKDIG
		ļ		l	Į.	AIARVHAENGELVAEGAKEAL
1				ŀ		DLGITGPEGIEISRPEELEAEATH
				ł		RVITIANRTHCPIYLVNVSSISA
				ł		GDVIAAAKMQGKVVLAETTTA
ĺ						HATLTGLHYYHQDWSHAAAY
		1				VTVPPLRLGIPPH
15399	45767	Α	15489	307	600	VSRRASSGSPVPLGSLHLQHSA
		1		•		PA/PSAAPKPRSPPARLSRPHLR
		l				GESAAPSPLSPGPASLAAAAAA
1	1					RVPAQPSPAPSTTPAPRTTLAPR
						VPHRLAPAP
15400	45768	Α	15490	3	222	
15401	45769	В	15491	1	1131	
15402	45770	Α	15492	136	201	
15403	45771	Α	15493	3	556	ELPRRLVCSKLRADPGRLTPDA
1				ì		CARPGMSRYLLPLSALGTVAG
1						AAVLLKDYVTGGACPSKATIPG
1						KTVIVTGANTGIGKQTALELAR
				ŀ		RGGNIILACRDMEKCEAAAK\D
		1		l		IRGETLNHHVNARHLDLASLKS
	İ		1			IREFAAKIIEEEERVDILINNAGV
		l		ŀ		MRCPHWTTEDGFEMQFGVNHL
		_				GHFLLTTWC
15404	45772	Α	15494	11	559	
15405	45773	A	15495	1	312	MAPAADREGYWGPTTSTLDW
		1				CEENYSVTWYIAEF\SWLMSGF
		1				LPTPSSLRDLTASRWVRSLPPSR
		1		1		SPAGRQPGPAEELPKASPCPWG
	L			L	Í	KSLSRPFASFSASSGPS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15406	45774	A	15496	2	1605	WQLPHPPAARSARQILRACQLS SVTAVAQSCLYGKQLC/GLTTG NIAGAGLLFFGGGIGGTILYAK WASHFRERVEKTIPYSDKLFEV LGSAPYNVPLPKKSTOSEPV LGSAPYNVPLPKKSTOSEPV LGSAPYNVPLPKKSTOSEPV LGSAPYNVPLPKSTOSEPV LGSAPYNVPLPKSTOSEPV LGSAPYNVPLPKSTOSEPV LSSVSEVTKEFKQPASOLOKOK GDTTPASATAGDTVLVPGVQHE ESLKTDHPAIDEGKFTPALSEEA SSSSIRERCEEIAACLAQEEKQ EQWKTESLAKSFEDALRGTANV TLQAIAAQNITVVQAVNAHSNIL KAAMDNSEIEGKKKSAKWHTV TLQAIAAQNITVVQAVNAHSNIL KAAMDNSEIEGKKKSAKWHTV TLQAIAAQNITVVQAVNAHSNIL KAAMDNSEIEGKKKSAKWHTV TLQAIAQODFFKRQLDSITPEVLPGW KGMSVSDPADKLSTDDLNSLIA RAHRHIDQLNRELAEQKATEK QHITLALEKQTLEEKVOEQELK YEFEQNLSEKLSEQELQFHHLS GOVINFTILDINTAVARLRGIE HAVQSIAVAEEEARKAYQLW LSVEALKYSMKTSSAEMPAVPL GSAVEALKYSMKTSSAEMPAVPL GSAVEALKANCSDNEFTKALAT AIPPESLTHGVYSEETLRVFFVA VOKLARRVAMID

	lon, /=possible nucleotide possible nucleotide insertion)
15407 45775 A 15497 89 2490 PKCCFF KVVLRI TGKIAG AKWDS	CLISGETA A AOSCI CG
KVVLRI TGKIAG AKWDS	CLESGETA A AOSCUCO
KVVLRI TGKIAG AKWDS	
TGKIAG AKWDS	PLRPCRRYSTSGSSGLT
AKWDS	AGLLFVGGGIGGTILY
	HFRESVEKTIPYSDKLF
1 1 1 1 1 1	PAAYNVPLPKKSIQSGP
	EVM\KESNRPASQ\LQK
	PASATAPTEAAQIISAA
	PAPAVQPEESLKTDHP
	PTPALSEEASSSSIRERP
	RLAQQEKQEQVKIESL
	DALRQTASVTLQAIAAQ
	AVNAHSNILKAAMDNS
	KSAQWRTVEGALKER
	EAADALLKAK*LLMDF
	EKMKSVIENAKKKEVA
	TAAEGKLHNMIVDLD
	VQAAQSEAKVVSQYH
ELVVQ/	ARDDFKRELDSITPEVL
PGWKG	MSVSDLADKLSTDDLN
SLIAHA	HRRIDQLNRELAEQKA
TEKQHI	TLAL\EKQKL\EEKRAF\
DSAVA	KAFRNIHRKLKYRA*TO
QKR*KE	EVRDAME\NEMRTQLR
RQAAA	HTDHLRDVLRVQEQEL
KSEFEQ	NLSEKLSEQELQFRRLS
QEQVDI	NFT\LDINTAYARLRGIE
QAVQSI	HAVAEEEARKAHQLW
LSVEAL	KYSMKTSSAETTS\IPL
GSAV\E	AI\KANCSDNEFTQALT
AAIP\PE	SLTRGVVTVKRPLGAR
F\YAVO	KL\ARRVAMIDETRNSL
YOYFLS	SYLQSLLLFPPQQLKPPP
	OALVEFEDVLGACNAV
	NQIYIAGHPAFVNYSTS
	GDSDDSRSVNSVLLFTI
	TTPTRLNVFKNDQDT
	PNLSGQGDPGSNPNKR
	LGDHPAEYGGPHGGY
	DEGYGPPPPHYEGRRM
	CYGPHADSPVIMVYGL
	NCDRVFNVFCLYGNVE
	KKQSPGGRPMGEEWL\
	DRAITHLNNNFMFGQK
	GAQAREGSRGTGERK
	PAEEHSEAEVLTHTEM
	SKQPAIMPGQSYGLEDG
	FSESRNNRFSTPEQAA
	IPSNVLHFFNAPLEVTE
	CDELGVKRPSSVKVFSG
	SGLLEWESKSDALETL
GFLNHY	/QMKNPSINLVT

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15409	45777	Α	15499	1	645	
15410	45778	Α	15500	3	677	SPPNPRAPYLFPASLQNPANWS
						PHFQYLPILSQPRSVLTAFRQQA
				1		QPRISMPQAQTPFRPQMERLDQ
				l		NIRRSNQPQHSLKPKLIQSKALT
				ŀ	ŀ	LFNSVKAEKDEEAAEEKFEVTR
				l		VVHIRGLIDGVVEADIVEALQE
						FGLISYVMVMSKKRQALVEFE
						DVLEACNAVNYTADNQIHIAG
				l		HPAFVNYSTSQKISRPGNSDD\R
				l		SVNSVLLFTILNPIYSITTDVLYT
15411	45779	A	15501	2	1712	DEQRRRSGAMVKMAAAGGGG
				l		GGGRYYGGGSEGGRAPKRLKT
	l		l	İ		DNAGDQHGGGGGGGGAGAA
				l		GGGGGENYDDPHKTPASPVV
1				l	c .	HIRGLIDGVVEADLVEALQEFG
			İ			PISYVVVMPKKRQ\SLV*FEDVL
1						GAGNAV\NYAADNQIYIAGHPA
1				l		\FVNYSTSQKISRPGDSDDSR\SV
				1		N\SVLLFTILNPIYSITTDVLYTI\
				1		CNPCGPVQRIVIFRKNGVQAMV
						EFDSVQSAQRAKASLNGADIYS
						GCCTLKIEYAKPTRLNVFKNDQ
						DTWDYTNP\NLSGQGDPGSNPN
						KRQRQPPLLGDHPAEYGGPHG
						GYHSHYHDEGYGPPPPHYEGR
						RMGPPVGGHRRGPSRYGPQYG
						HPPPPPPPPPPEYGPHADSPVLMV
				į		YGLDQSKMNGDRVFNVFCLYG
			İ			NVEKVKFMKSKPGAAMVEMA
			İ			DGYAVDRAITHLNNNFMFGQK
		ŀ				LNVCVSKQPAIMPGQSYGLED GSCSYKDFSESRNNRFSTPEOA
						AKNRIQHPSNVLHFFNAPLEVT
1		1				EENFFEICDELGVKRPSSVKVFS
1			1			GKSERSSSGLLEWESKSDALET
	1		i			LGFLNHYOMKNPNGPYPYTLK
15412	45780	A	15502	 	528	NHOIRNDFTISPGVKADSRTSPI
13412	43/60	^	13302	ľ	328	POOPASSEDITEAAVSEAKDSLA
1				i		GGVAAAISKMAVAPIERVK/RV
1						PKEHGVLSWCGNLASVIRYFP\
1						TOALNETEKDKYKOIFLDGVDK
1		l	1	I		RSOFWRYFAGNLASGGATGAT
		l		l		SLCFVYPLDFAHTRVAADVGK
			1	l		AGAERELRGFGDCLVKIYKSDG
			1	l		IK
15413	45781	A	15503	1	1452	
	1.270			Ľ		

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15414	45782	A	15504	2	991	RSAGVKAGSRSPVLQGSASSF NMTDAAVSFAKDFLAGGVAA AISKTAVAPIERVIKLLUQVQIH ASKQIAADKQYKGIIGCVVRIP CQALNFAFKDKYKQIFLGGVDK RTVGFWRYFARNLASGGAAGA TSLCFVYPLDFARTELAADVGK AGAEREFRGLGDCLVKIYKSDG IKGLYQGFNVSVQGIIYRAAYF GYYDTAKGMLPPPKINTHIVIS WMIAQTVHCCCPG-LPYPFDT RSV/RRNEWMGSRKIGTDIMY TGTLIDCWRKIARDEGGKAFFK GAWSNVLRGMGGAFVLVLVE YSKKYT
15415	45783	Α	15505	1	613	
15416	45784	Α	15506	1	1695	
15417	45785	Α	15507	131	723	LLEGKLTNKDIHTKNPSVRIV RSSKTKER VSAMEDEINEMKRE EKFREKRYKRNEOSLOEI WDYL KRPNLRLIGVPESDOENOTKLE NTLQDIIQENFPNLAROANIQIQ EIQRMPQRYSLRRATPRHIVRF TKVEMKEKMLRAAREKGRVT HKGKPIRLTADLSAETLOARRE WGPIFNILKEKNFQPRISYPAKL
15418	45786	Α	15508	1	3156	
15419	45787	Α	15509	1	879	
15420	45788	Α	15510	1	804	
15421	45789	В	15511	1	1230	
15422	45790	A	15512	2	827	
15423	45791	Α	15513	1	1011	
15424	45792	Α	15514	1	1722	
15425	45793	В	15515	1	780	
15426	45794	Λ	15516	1	2712	
15427	45795	В	15517	1	855	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15428	45796	A	15518	3	1263	GRTIOTKGKEVENFERNLEECI TRITNTEKCLKELMELKTKARE VREECRSLRSQCDQLEERVSAM EDEMNEMKQEGKFREKRIKIN EGSLQEIWDYKRPINLEIU EGSLQEIWDYKRPINLEIU EGSLQEIWDYKRPINLEIU ENDARGANVQIQERQRTPQRYSS RRATPRHIIVRFTKVEMKEKMG LLVPNWTNHSPLERALIFDYKG FCRECTTHGJIGSPAGANOROP LAATLSCPGGEGQSAVARLTGL IGVPESDVEKETKLETTLQDIIG ENFPILARQANVQIQEIQRTPOR YSSRRATPRHIIVRFTKVEMKE KMLRAAREKGRVTLKGKPIRL TADLSAETLQARREWOPIENIL KEKNPQPRISYPAKLSFISEGEIK KEKNPQPRISYPAKLSFISEGEIK STOKOMLROFVTTRRALKELL
						KEALNMERNNRYQPLQNHAK
15429	45797	A	15519	1	1137	
15430	45798	A	15520	1	1578	
15431	45799	Α	15521	1	1656	
15432	45800	Α	15522	1	1120	
15433	45801	В	15523	1	1541	
15434	45802	В	15524	1	1632	
15435	45803	С	15525	53	352	
15436	45804	A	15526		973	MGKKQNRKTONSKMQSASPPO KERSSSPATEQSWMENDFDELR EGGFRENYSELREDIOTICKE VENFEKNLEECITRITNTEKCLK ELMELKTKARELQRVSAMEDE MNEMKREGKFREKRIKNEQS LOEIWDYVKRPINLRIGVPESD GENGTKLENTLQDIDGENFPNL ARQANIQIQEILRIPQRYSSRA TRIHIVRFTKVEMKEKMLRAA REKGRVTLKGFIRLTVDLSAE TLQARREWOFIPNILKEKNEP RISVPAKLSFISEGEIKYFTDKQ MLRDFVVTTRPALQELLKEALN MERNNEYQPLQNHAKM
15437	45805	Α	15527	1	2896	
15438	45806	В	15528	50	658	
15439	45807	А	15529	1065	1260	RSHALALDSAGSSSPESPH*RAS IPHTALGQNAGSWAGTAHSHH GPFLWSDPRHPQVVPHRTCCP
15440	45808	Α	15530	96	219	EFSTKPRLSGP*SSWEMLILEPL APDTTIDLKLLYSDRDF

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
		<u> </u>				
15441	45809	Α	15531	5459	6324	LLANVSQLLADISQLLANFTQL
			İ			FSHFSQLLTYLSKLFTHLPQLLT
						HFPKLLTHQPELFSNQSQLHPNI
		1				T\RYSPTSPSYSPTSPNYTPTSPN
		l				YSPTSPSYSPTSPSYS/RDLTKLL
						PFQPTIHTTVSNLYPKLTQLQPQ
						FAQLQPNLTQV\PPTSPSYSPSSP
		1				EYTPTSPKYSPTSPKYSPTSPKY
1		1				SPTSPTYSPTTPKYSPTSPTYS\Q
						PLQSTPQPLPSTHLLAPLTRPLP
				İ	1	PSTRPPAPPTRPPPPKAQPTLPLP
1	1	1				LVTRPPAPPTVSQARLSARMTV
						TRRTEGTWGAAAG
15442	45810	Α	15532	13	941	SVVLRPLLVVAAATATTPVAA
		1			ł	AAATNTAKKTATVMVGEMVA
		1				GERGVLQMGHDITRFFYSNIQT
					l	VINNWLLIEGHTIGIGDSIADSK
			ļ			TYQDIQNTIKKAKQDVIETFEN
	İ					QVPRILNDARDKTGSSAQKSLS
1	l					EYNNFKSMVVSGAKGSKINISQ
		l				GGGSQGGRGGPPISMWLLALC
1		ļ				LVGLAGAQRGGGGGGGAPG\
1		l				GPGLGLGSLGEERFPVVNTAYG
		ł				RVRGVRRELNNEILGPVVQFLG
		l			}	VPYATPPLGARRFQPPEAPASW
ł		1				PGVRNATTLPPACPQNLHGALP
		1				AIMLPVWFTDNLEAAATYGGC
						FRVPKKA
15443	45811	Α	15533	207	696	TNLQEKNKQPHQKVGKRYKQT
				ŀ		LLKRRHLCSQQAHEKMLIITGH
		i i				QRNANQNHNEIPSHTS*NGDH*
		1				KVRKQQMLERMWRNRNAFTL
ı		1				WVGV*TSSTIVEDSMAIPQGSR
		1				TRNTI*LSHPITGYIPKGL*IMLL
		1				*RHMHTYAYCGSIHNSKDLEPT
		L				QRSINDRLD
15444	45812	Α	15534	1	307	
15445	45813	A	15535	1	381	LPPPHDRLWCVMFPTLCPSVLI
		1				VQFPPMSENMRCLVFCPCNSLL
						RMMVSSFIHVPTKDMNSGIL*S
	1	1	1			RPADIELFSINLVNKKIFLKGKF
	1	1				GDPLSLQRQLRNIVFSMPCCFPS
		L-				FLLPWVFSQLSVKPH
15446	45814	A	15536	134	278	DPAFLAPSPV/LMKQAPQATSG
		1				LMEPPGPSTPIVQRPRILLVIDD
	15015	<u>_</u>	15507		1000	AHTD
15447	45815 45816	B C	15537	1	840 891	
13448	43010	<u></u>	13338	11.	091	L

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino aeid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15449	45817	Α	15539	1	3102	MYSHVITVCRHVKNGDILLLNR
						QPTLHRPSIQAHRARILPEEKVL
	ĺ			l		RLHYANCKAYNADFDGDEMN
		ĺ				AHFPQSELGRAEAYVLACTDQ
			ŀ			QYLVPKDGQPLAGLIQDHMVS
						GASMTTRGCFFTREHYMELVY
						RGLTDKVGRVKLLSPSILKPFPL
İ						WTGKQHINRTKDKNHMIISIDA
						EKAFDKIQQPFMLKTLNKLGID
i		1				GTYLKIIRAIYDKPTASIILNGQK
1	١.	1				LEAFPLKTGTRQGCPLSPLLFNI
		1			1	VLEVLARAIRQEK
15450	45818	Α	15540	1	779	IPMKDHDAIKLFIGQIPRNLDEK
		1			ŀ	DLKPLFEEFGKIYELTVLKDRFT
		l				GMHKGCAFLTYCERESALKAQ
i						SALHEQKTLPGMNRPIQVKPAD
		l				SESRGEDKKLFVGMLNKQQSE
		İ		l		DDVRR\LFEAFGNIEECTILRGP
				ŀ		ERQTAKGCAFVKYSS\HAEAAG
						RPSTRLHGSQTMPGAFVPVWV
						GQVSADTDKERTMR\RMQQMA
						GQMGMFNPMAIPFGAYGAYA
	i		1			QALMQQQAALMASVAQGGYL
			İ			NPMAAFAAAQMQQMAALNM
						NGLAA
15451	45819	Α	15541	1	1413	MEYYAAIKKDEFISFVGTWMK
				Ì		LETIILSKLSQMQKTKHRIFSLIV
				ļ		LCDITDILKEFTVAEAGGFALNE
						IMKEICKSRFSAKVSLHWKMW
	1					HSTASLGYHRVNVTNPDPLNLC
						LGHCWLGLRDTGQKQACGGP GVCPGMFRRACGGNAPCKEAP
						SVTEASSPDGSVLCLGSHVPITP
		l				APLSSAQMVCHMVEDRRTVEL
ł		ľ				ILGATHNLFFSTSPLTWSLRAIP
						GNCVLFLKDYVTEPVRSQSILHI
1						AGPTFKOTPNLPFLKFPEATKA
				1		HGPNVYQNDSTCPEPTAQVSH
1	1	1				WSSVSSTLIASPVPGPEDRKLFV
1		l				GMLNKQQSEDDVRRLFEAFGN
						IEECTILRGPDGNSKARPLASSQ
1				1		GAPSVKYSSHAEAQAAINALH
						GSQTMPLMQQQAALMASVAQ
1		1				GGYLNPMAAFAAAQMQQMAA
	1					LN\PNGLAAAPMTPT\QANGQP
1	1					AAEENFANGIHPYPAQSPTAAD
						PLQQAYAGVQQYAGR
		Ц_	L	L	L	L 244 ING 144 INGK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15452	45820	A	15542	254	883	DLRKDSPWEHRQCDKREPGNS APPAPHSWGFAVFNRAGDVR QLCKRADTRITSPRAPFGSAGA CRGCGRAGARSLASPKARTCIS SARRRAGGPULGAGSGGAKR PDR*TRREGANA*KTI.HI./HSSL APSPGGVLAGFQEPPGRPQVAC CAATAAGGIRAGERRPAAAAA AAPWAQKLERLRLQRRGPPGC R**MYSPRELFSGSTSR
15453	45821	А	15543	1	768	MCDFGKSTSLPASPTSSSANGR DNSTYTRLEGGGFMISVASRO SCAESARYTDAHVAKSGYGAV TPSSYGANI.A ASLLEKEKLGFK PYPTSSFLTRPRTYGPSSLLDYD RGRPLLRPDITGGGKRAESQTR GTERPLGSGLSGGSGFPYGVTN NCLSYLPINA YDGGVTLTQKLD SQSDLARDFSSLRTSDSYRIDPR NLGRSPMLARTRKELCTLQGL YQTASCPEYLDVLENYGKG SASQVPSQAPPSRVP

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15454	45822	A	15544	85	2249	SAEAMSGDGATEQAAEYVPEK
		ļ				VKKAEKKLEENPYDLDAWSILI
		1			i	REAQNQPIDKARKTYERLVAQI
						PSSGRFWKLYIEAEIKAKNYDK
					ŀ	GEKLFQRCLMKVLHIDLWKCY
ĺ		1				LSYVRETKG\KLPSYKGKMAQ
		1			l	AYDFALDKIGMEIMSYQIWVD
		1			l	YINFLKGVEAVGSYAENQRITA
						VRRVYQRGCVNPMINIEQLWR
						DYNKYEEGINIHLAKKMIEDRS
		İ				RDYMNARRVAKEYETVMKGL
						DRNAPSVPPQNTPQEAQQVDM
1						WKKYIQWEKSNPLRTEDQTLIT
		l			i	KRVMFAYEQCLLVLGHHPDIW
l						YEAAQYLEQSSKLLAEKGDMN
			ĺ			NAKLFSDEAANIYERAISTLLK
						KNMLLYFAYADYEESRMKYEF
	1					VHSIYNRLLAIEDIDPTLVYIQY
	İ]			MKFARRAEGIKSGRMIFKKARE
						DTRTRHHVYVTAALMEYYCSK
l						DKSVAFKIFELGLKKYGDIPEY
						VLAYIDYLSHLNEDNNTRVLFE
						RVLTSGSLPPEKSGEIWARFLAF
		İ		ł		ESNIGDLASILKVEKRRFTAFKE
						EYEGKETALLVDRYKFMDLYP
		l				CSASELKALGYKDVSRAKLAA!
		l			1	IPDPVVAPSIVPVLKDEVDRKPE
		l				YPKPDTQQMIPFQPRHLAPPGL
		l			i	HPVPGGVFPVPPAAVVLMKLLI
						PPICFQGPFVQVDELMEIFRRCK
		1				IPNTVEEAVRIITGGAPELAVEG
		<u></u>				NGPVESNAVLTKAVKRPNEDS
15455	45823	A	15545	12	427	SAEAMSGDGATEQAAEYVPEK
		1				VKKAEKKLEENPYDLDAWSI\L
			i			IREAQV**YRITFPSLWVQSHYL
		1				VLPVNNIFIILTL*MLFIGFQF*N
						QP\IDKARKTYERLVAQFPSSGR
		1				FWKLYIEAEVTILFYFFLYQYCS
						IHL
15456	45824	A	15546	150	469	SAEAMSGDGATEQAAEYVPEK
					Ī	VKKAEKE\LEENPYDLDAWSIL
				l	l	REAQNQPIDKARKTYERLVAQF
l	1			1	I	PSSGRFWKLYIEAEVTILFYFIFS
		L				YISIAAFTVVIENKLGT
15457	45825	Α	15547	1	834	
15458	45826	Α	15548	443	732	SFMRWQKGQLPQLCLHRESHC
				1	1	SPYPQPSFLGMEP/ADIHETTFN
l	1		1		1	SIMKCDVDIHKGLYANAVLSSC
				1		TTMYPGIA\NRMQKEITALAPS
l						MMKIKIIAPA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ŀ				sequence		
15459	45827	A	15549	1	1214	MEEEIAALVIDNGSGMCKAGF
10.00	15027	`		ľ		AGDDAPRAVFPSIVGRPRHQGV
l						MVGMGQKDSYVGDEAQSKRG
1		ı				ILTLKYPIEHGIVTNWDDMEKI
		1				WHHTFYNELRVAPEEHPVLLTE
						APLNPKANREKMTQIMFETFNT
						PAMYVAIQAVLSLYASGRTTGI
	į.					VMDSGDGVTHTVPIYEG\YALP
	ŀ					HAILRLDLAGRDLTDYLMKILT
l	l			l .		ERGYSFTTTAEREIVRDIKEKLC
						YVALDFEQE\MATAA\SSSSLEK
						SYELPDSPRSSTNWQLSGFRLPL
						RHSFQPSFLG\MESCG\IHETTF\
				ļ		NSIMK\CDVDIRKDLY\ANTVLS
	İ					GRTPPCTLGIADRMQKEITALA
		ļ.		1		PSTMKIKIIAPPERKYSVWIGGF
		1		ĺ		HPWAWLSHLSSKMWVQQARS
1	}					MTESGPLHSSTANGFLGGLLTL
	l			1		VAFTPFLWTKP
15460	45828	A	15550	1	2355	MKQITFAPRNHLLTNTNTWTPD
13400	43020	n	15550	l'	2333	SQWLVFDMRTSGASFTGETIER
				ŀ		VNIPTGYVEKPNKOCRHSVTRY
		1		İ		GHGGHSDHTPSLTVATVATQIT
						PRHSGHGSHCDRNLSLATATA
						ATLITPHHLLWPRWPLGSHSVT
						WCGHGGHSDHTPSPTTATAVT
		1				RITSRHPLRPQQSLGSHPVTHYS
						HGSHSDCIPVTRYAHGGHSDHT
						PSLAMATAATQMALCHSLWPR
		1				OPLGYLVNKIRRKRCTDGVSPS
	1	1		1		GKAPVFDTGIPWFESRYPSHLL
	1			ł		RVSGSPPGYWGIAKRHQILNVS
				ŀ		NMFGKFKTNLLGYRQAVRHRI
				1		LIPAFRGSNPRTPANLFKTLTLIS
						NMRALENDFFNSPPRKTVOFG
				1		GTVTEVLLKYKTAIEKRLEKLA
				ŀ		VDPHRDRAQMLKVENVQQAW
						QQWINKLPPARREDEDVKEIR
		1				WMIEELRVSYFAQQLGKNSRA
		1		1		ESMLWSFHKAAIEAARTNIIQA
	i	ı				QTRVEAAQATERRIAADIDDSE
				1		LKAPRDGRVQYRVAKPGEVLA
			1			AGGRVLNMVDLSDVYMTFFLP
						TEOAGTLKLGGEARLILDAAPD
	1	1		1		LRIPATISFVASVAQFTPKTVET
		1		1		SDERLKLMFRVKARIPPELLQQ
		1		1		HLEYVKTGLPGVAWSFLYSINO
		1		1		TICLRLDSIEAKLQALEATCKSL
	1	1		1		EEKLDLVTNKQHSPIQVPMVA
				l		GSPLRTTOMCNKVR/CVNP*AT
		1		1		VVPPPVPQPTTQQYQGLDAGA
	1		L	<u> </u>	l	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15461	45829	Α	15551	1	956	MSSGRRDALARGLGNSGDGAP
				[HLGEHROLEDVGGVTVTVTWT
						GRLOAAGCWTRSVSSTAAAGR
						REPSTRDGAALSTEPPDPPERLG
						HRGSRAGLTALGHVRTRGSSO
						RGGRPEPWEHWLENGTRRTEL
		ĺ			0	VDGVSYESWRRPCRGOLESR
						LVLENHVATDEDEPALKCORL
						EINCODPSIKSFLYSINOTICLRL
						DSIEAKLQALEATCKSLEEKLD
						LVTNKOHSPIOVPMVAGSPLRT
						TOMCNKVR/CAMHWLCVSCIF
						TCLPGWRPAAPDQGPAAISLCS
		ľ				LPSSSOGHREPLALGLPSALPPA
		ŀ				HRQRLRGSATVPIPP
15462	45830	В	15552	161	925	IIIQIGIGOSAI VI II I
	45831	Ã	15553	31	856	GGFLHHASFFVGRGRTLLMYFS
	15051		1.5555	· ·	***	LLGATPKPEPEPEQVIKNYTEEL
					İ	KVPPDEDCIICMEKLSAASGYS
						DVTDSKAVGPLAVGCLTKCSH
·						AFHLLCLLAMYCNGNKGPEHP
. I						NPGKPFTARGFP\ASATFQTTP\
						GPQELVPYTPQITAWDLEGKVT
						ATTFSLEOPRCVFDGLASTNDT
			İ			VWLVVAFSNASRGFQNPETLA
		l				DIPASPOLLTDGHYMTLPVSPD
		l				OLPCDDPMAGSGGAPVLRVGH
. 1		1				DHGCHOOPFCNAPLPGPGPYR
. 1		ı				WVVPTGALGLGLCLG
15464	45832	Α	15554	827	1371	GSSAGEASSSLRRLSPSOPISRPL
.		1				LPAGEKVLVVWIEDOTSHNIPL
. 1						SQOSKT/TLFNSMKAERGKEVT
ı		1				EEKLEARRGWFMRFKERSCFPN
		l				VKVQGEAANADIEAVACYPEY
		l	1			LAKITDEGGYTEQQIFSVDETA
		l				LYWKKMPSRTFLARRKNCHSH
		İ				PSFSSQHLVQSAAINIKAKSSTS
						KKIMTP
15465	45833	Α	15555	799	906	FLNKLAFALRCGLALNSFLRKI
1						QEPSLG\SKSGSLS
15466	45834	Α	15556	143	560	

****	WO 01/0/5067 PC 1/0501/08051							
SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,		
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)		
				sequence				
15467	45835	Α	15557	1059	1940	QQMQQEMRRKSKWMAMLGK		
						WETYQNS/TKGHLMKRVYKGI		
1				ŀ		PMNVRAEVWSVLLNIQEIKAK		
1						NPRQYRVMKEKGKRSSEHIHQI		
					ĺ	DLAVSQTLRNHMFFRDRYGTK		
						QQELFYILLAYSKYNPVVGYCR		
						DLSHIAALFLLYLPEEDTFWAL		
						VQLLASERHSCRVGEQLPLGLH		
					i	AARPGDGHPGOPRHLPCVASLL		
						GAFRTSLLTVPKEPGTDPPEPKS		
		l				DAFHPHQQKASHPPRGCPLCPG		
		l				ATPSGSDSVQLALRNLRVLLPF		
		1				SCSGSSCPWYPRKEALHPGGLE		
		l				EQRGLQPLMGLVTG		
15468	45836	A	15558	123	448	ALAAAVYPSHSERTGEEAKKIC		
						SVR*LLRSAQEEI/FLRGKNSRT		
		l				GVL*KLLORK*RYORALGLAM		
						LSEWTISPKGPVSTGITKPMLTA		
		1				SRVTVPRMTMSALLDFIHGA		
15469	45837	Α	15559	1	6387			
15470	45838	Α	15560	52	3218			
15471	45839	Α	15561	3602	4015	GSVESPLGQKTPRTSSASPCWPS		
		ı		i		TNLYRMELELYGPHVTSQVVK		
		l				STRCAPIRWSAAPSILCEVRVPV		
1		i				TMWTCLPLPATGRLSMW*IWP		
		l				RQWPCVLTSATQS*LTRCGGGT		
1		1				RAVSLAPQSHL*VCPAQSSWTS		
1		1		l		IILWT		
15472	45840	Α	15562	932	1316	RVQDIPAKRGIPQFRPKIKGIGQ		
		1				SYKQHYPARPWNVLAQKVV/V		
1		l				FEDQDALLPCLLTDPVLEAGVS		
1		1				LVRVRGRPLMRHTNYSFSPWH		
1		l				GFTIHRAKFIQSQDYQCSALMG		
		_				GRKVMSISIRLKVQKAVNRA		
15473	4584I	Α	15563	I	263			
15474	45842	A_	15564	66	185			
15475	45843	Α	15565	I	4217	MRLAEERAALAAENADGEPGA		
	l	l			1	DRRLRLLGTYVAMSLRPAAGA		
		l				WERCAGSAEAEQLLQAFLGRD		
		l				AAEGPRPLLVVRPGPRGLAIRP		
1		I		l	1	GLEVGPESGLAGAKALFFLRTG		
1	-	l				PEPPGPDSFRGAVVCGDLPAAP		
1		1		l		LEHLAALFSEVVLPVLANEKNR		
1		I		l	l	LNWPHMICEDVRRHAHSLQCD		
		l		l		LSVILEQVKGKTLLPLPAGSEK		
		1		l		MEFADSKSETVLDSIDKSVIYAI		
		ı		1	ŀ	ESAVIKWSYQVQVVLKRESSQP		
		L_				LLQGENPTPK\VELEF		

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15476	45844	A	15566	26	494	RPGCGEWLGNAELWGQECQL HYSPPVSPWENSSGPRFSISKTG SLIPPLLNDICQNQKPQIILSHPA RCLGLLDHFPQQWERRQAONR AETG*GQRSAQGQTFRQRPLSV SGNCRVPPRATVSRK/PRPAPA Y ALSLKSPRRPPRTGPAPPGLYP
15477	45845	Α	15567	1	1242	
15478	45846	A	15568	1114	1219	VDLPPGAPVRQSGSHGSVLRDH KTEGEARVLPLHQRVGKAGGR GATTPGHVHLEPDVTICCGHSEC RPIC/ICLSWEEAQSPANCRACR EPSQKKDFKTNILLKNLVTIAR KASLWQFLSSEKQIGGTHRQTK KMFCDMDKSLLCLLCSNSQEH GAHKHYPIEEAAEEHREKLLKQ MRILWKKIQENQRDLVEERRT AFLWRKRTLRLKGMTSGAQGT SQKVALGLLQQPLCHTVRAQS NSARVHKQRGGDAVSSGAGIP DPKAQRFPATLRSQAFTCTAGR RWGSGETVLHFLQRAGVRGAV AELHRQLLLSSSAAGFECEGL CTSGPHITQGPDHQELYLSHEG

1982

						PC1/US01/08631
SEQ ID			SEQ ID NO:		Nucleotide location of last	
NO:	of peptide sequence	hod	in USSN 09/540.217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		03/010,21/	sequence	or popular conjunct	
		<u></u>				<u></u>
15479	45847	A	15569	1	2694	MRMAATA WAGLQGPPLPTLCP
						AVRTGLYCRDQAHAERATDVV
	ł					LLAPFCQPKTRSHGTCPPTERDP
	ł					RGEGSTEYPGRVDGIQGWGTR
					1	ALTGWTDRRLLCQACQTLPPR
						HWFLPGARGWLGGSPCAHGQE
						SLPSQPSPILLRVESVKSRMLRR
ŀ		l				RAQEEDSTVLIDVSPPEAEKRG
						SYGSTAHASEPGGQQAAACRA
ł		l				GSPAKPRIADFVLVWEEDLKLD
						RQQDSAARDRTDMHRTWRETF
						LDNLRAAGLCVDQQDVQDGN
						TTVHYALLSASWAVLCYYAED
						LRLKLPLQ\DYPTRPPTGRPAC/
						CAWLGIPNVLLEVVPDVPPEYY
						SCRFRVNKLPRFLGSDNQDTFF
						TSTKRHQILFEILAKTPYGHEKK
						NLLGIHQLLAEGVLSAAFPLHD
						GPFKTPPEGPQAPRLNQRQVLF
		l				QHWARWGKWNKYQPLDHVR
1						RYFGEKVALYFAWLGFYTGWL
						LPAAVVGTLVFLVGCFLVFSDI
						PTQELCGSKDSFEMCPLCLDCP
		1				FWLLSSACALAQVREEAGRLF
		l				DHGGTVFFSLFMALWAVLLLE
		1				YWKRKSATLAYRWDCSDYED
		l				TEERPRPQFAASAPMTAPNPITG
		ļ				EDEPYFPERSRARRMLAGSVVI
		l				VVMVAVVVMCLVSIILYRAIM
l		1				AIVVSRSGNTLLAAWASRIASL
		1				TGSVVNLVFILILSKIYVSLAHV
		_				LTRWEMHRTQTKFEDAFTLKV
15480	45848	Α	15570	1	924	MISQFIPENYFSGKSSATSGWA
						RAQRDFRVGAGSARPALLSGRL
				ĺ		APPALGSERLSTRASSCGGLPR
		l				ATPRGRAPPPATRHPVPSTAQR
		1				LRSAAARQRLAGSSARGRDAL
		1				GKASWAPESVLEVLARAVRQE
		1				KEIKGILLGKEEVKLSLFAGDMI
		1				VYLENPIISAQNLLKLMSNFSK
		1			1	VSGYKISVQKSHAFVYTNNRQS
ĺ		l		1		ESQIMSELPFTVATKRIKYLGIQ
	1					LTRDVKDLFKENYKPLLNEIQE
		l		1		DTNKWKNIPCSWVGRINIVKM
ı						AILPKVIYRFNAIPIKLPMTFFTE
	122.10	I.—	1.000		004	LEKTK\FIWNQK\RAHIAKTIL
15481	45849	A	15571	1	825	
15482	45850	Α_	15572	1	1488	

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				,		
15483	4585 I	Α	15573	I	1476	MPIKVDWIKKSRTHTPRDTMSF
	1	1				KRGINISIQEYRNTPDASILSELR
						LERDTNTQVYTYEQEHTNKNN
l		1				GHQAIDKQNKGHIRRESDRGEY
						YPLDCRPTGSLTDKGWGMLLC
						GMMTAGKPTPILTFLLDVGLPG
	l					GTAINKSSVIRVRNREEGNVEK
						WGERQVDQRDAVMRVRCGIR
						NNVEAGLKYGPGTMKVSLSVF
						YTTQQCRRYLSVLCFAGLARAS
		ł				VPFNEDFQMSLDEKLFSNLHPL
1		ļ.				GSSSSPPKSLLHCPLLEHQFTML
						GFSDLGAQLASPSGSHTRAAGV
		1				AACQPAPTLTWLSPWVARGLG
	1	l				TLEQGVVLVGEARAAQEPMEW
						VGGSCMVGCRSRALPRGKAAK
						AWREIQRSAGAGGAPVSPWSG
						GAPSRACPPSASSVRFLCAARR
						LACNTYSPVLYCVRNWWILGL
	ļ.					TDFNNEATDPRDPFLANVPGQL
						AAVAIGTCDGNVYSAGDRDYR
				l		FALESILKACTLALALEEVGPPA
						VQDKIGADP\TGLPF
15484	45852	Α	15574	140	448	GFLGACAGREFKMAPP*SSLSG
		ŀ				*KLSVSPCPGLLISAEGPEAVFSI
		ļ.				V*NARVRGFILEVSETKNPPIPD
		ľ				TISTKHASLMSDSYREKKLVPV
						VYPHSSTPSFQ
15485	45853	Λ	15575	1	1001	MEAPQGALEQDGPAVLSQQRH
					1	RKKPFTATPIDPSYHRSHAISLS
				l		VIIGYCVPPSHAAEMAAYGWT
		ľ				MGKSKLMAGQWITQSQVFCDS
						STDGPRQHKAQTAILTKLQSQQ
						RSTNNSTVSHRQQWVNCRLLR
				ŀ		VKCQRLRAHKELLPVQRWLWR
						QGRCVPTTSLVAVSETAGAGA
			ŀ	1		AALALSRLCRCLQVSASARPFS
						SRTLSPQPKCWAMAASHQPIKG
İ	1	1	l	l		ILKNKTSRTSSMVALSEQPCRT
		1		1		VHEELSKKCQKWDEMNILSTY
	1	1	1			HPADKDYGLMKIDEPSPPYHG\
1	1	1	l			LTGDDENACSDTETTETMVSDI
		1	1	1		LAKKLAAAEGLEPKYWVQEQE
	15051	Ļ.	15506	ļ. ——	1057	SSGEEDLLAEE
15486	45854	A	15576	73	1851 418	CKBDCDKACOM I I MACAGET
15487	45855	A	15577	/3	418	GKPDGDKVSQHLLLWKG/YSF/
		1		I		PPSLMKLSLAGYEILS*KFFSLR
				1		TLNIGPPLSSGFRVSAERSTVSL
		1				MDFPLCITQCFSPAALNIFSFIST
15 400	45856	A	15578	2	460	LVNLTIMCLGVALLEQYLCCVL
15488	43836	Ι^	113378	12	400	<u> </u>

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
15489	45857	Ā	15579	1	2031	MATEVFLIPYALLOCSLASYPS
,						KPGIYVPSPQMWVGSVTCSDQ
						KNTIEVLVPKAYTLAROOYKSL
					l	SEQVYSCHQGTLNTRAQESAAS
						LLKPARPRTRQKKETPNTSEHQ
1		1				KEOTLDTPPLRTVTLTTRVRGFI
					1	LEVSETKNPPISDTMGLDISKOE
		ŀ				,
		ļ.			1	EVLGRESDEQGEQSQGLSEMQ
		ĺ				TALAMLLTREPISAKQAPHEPP
1			1	l		RRQEDNTQAREGETRSRKTRA
					l	EGQKGHAHAENERNAKERESE
						RKRQEAEDENGKEKQERNEGT
						HARHMKRRKRDEKKRQGQNK
					1	PGNKRPEGNEEAQRRRKKNRE
	1	ŀ			1	RQRFSFYQSGPSAAGLLVFAGG
					1	PLQSLFAWVSPLEAAEQQRFLS
						VPSSGPLRKFCLYFGFVGNATP
						LRGLNTHGRLLTPAHGTRRRHT
						RPLNSPVPEKAGSTTESNGSPY
						GONPGPLPPKGNRAOGEKWM
						GRRIQGTTRPRAREANHKKMT
1				İ		OIMFETFNTPAVYMAIQAMLSL
				1	i	YTYGCTTGIVMDSSDGVTHTVP
						IYEGYALPRAILCLDVAGRDLT
			i			NYLMKILTEHSYSFTTMAEREI
ì			1			VCDIKEKLYYVTLDFEQEMAT
				l		AASSSLEKSYELSDGOVITIGN
				[ELSCCPEALFQPSFLSMESCDIH
						KTTFNSIMKCDMDIRKDLYTNT
				l		VLFGGNNMYPGIANRVQKIVTL
		Ļ.				VWMGSSILALLSTFQQMWVSK\
15490	45858	Α	15580	485	886	VPYFHFTCSHSSASQLPFLQMM
		1		1		KIAKTCSLGLNRLPEAKLKAFP
	i		1	į.		TPRVTSL/WPRQRGRPRR\GCGR
				l		NRSGSLVSAARGVGGRSQAAA
				1		ARRDAPSPPATSRMPPRQGSVS
				1		WGCARNAPGRRADPEVRRQPG
						PSCT
15491	45859	Α	15581	1	435	
15492	45860	A	15582	2	339	HRCSTGAFSTAASNSNADRPPP
1		l		l	1	ARRRGLPR/RGPP/SRRRSPRPPR
1				1		REPWPRTRRPRTRTAKARRRAT
1				1	1	LEGRCRRACGHWQPRAAAAA
1					1	AAAAAAVAAAAAAAAAAA
1					I	GASATW
					<u> </u>	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15493	45861	Α	15583	1067	2093	RRNTSSKCRSRNCRCPRSPHRH
						RPSLRPRSQQPKCKCRPRSRRSS
ì						RAPSSRRSRPQGLVPC*RAPPW
						PTSRWPGSPGFPLLNCSRKGRC
		1	1			RPRHPSQPRCPCRSLRWCPSRQ
		l				LWSPHRESPPCP*TSRGSAWRS
	1					VSHRRQQDRPWWTQPVHMQQ
		1				LLKLKQQAVQQKAIQPQAAQ
						GPAAVQQKITAQQITTPGAQQK
		l				VAYAAQRALKNPVSYHTHLPG
		1				PETGRGPASADPDPGCKTSSSCS
l	1	ĺ				TANTRGQHPASCLCFPAGFSTD
1		i				CGAHAGDGGRAAGADDPCSDF
		1				DCPGGSAETHSAAGGDHGVGP
1		- 1				APDSRRSQPSPGARQLRQPKPA
1	1	İ				AQVTDEGPCCQAKDTY
15494	45862	Α	15584	10	607	RALCGESRLPLARKLS*AKKSV
						QRSTLQSRRIWVCWFHFAN*SL
ļ						F*KQGRT*ESPL*L*LIPAS*RPS
						TSESPPL*KANFQQPHRGL*EKV
1	1			1		AEGRRGP**EY\YTSSSSSSSSS
	1					ssssssssssssssssssssss
	i			l		SNNSSNSRHRRRRRRRLLSFLSF
						RRRHSRCCLSRSRPRRRPRRHP
l						ARLWLRSRCTDQRKNFQLPRK
						TV
15495	45863	Α	15585	217	1012	AKNDSKRONFGHRGQPTFGAD
	i					KAKGERVRTSSTIRRTSYLDTIT
	ļ					GPYLTGQWPRDPHGHYPSCMK
						DKATOTP\SCWAEEGAEKRSHQ
						RSA\SWGSADQLKEETPTDLOQ
	l			1		NAQQKLYKPEEVGCLISDFLKE
l						KENCOPRILYAGKLSSECTPDE
						EAVNIVEMITKDLEYYINLVDK
	i					EWQGLRGLTPILKEVLRVKCY
						QTASHATKKSGMKGKVSQFGK
						LHCYFKKLPOLPLPSATAILISO
						LTSTRQELHHKKDCNLLKTQMI
15496	45864	Α	15586	2	387	DIOTROPENING DO VEDETONIA
15497	45865	A	15587	1	165	LFMFFCLLYIR*OLLHPGYNAPF
1.545/	7,000	1	13367	I.	1.00	QNTAATRGRGIAQTAPEAAFQL
		1	1		1	FLHORVRFHC
15498	45866	A	15588	172	359	PDGDGIESINYLGQYGHFHDIDS
13498	4,7000	^	13308	11/2	337	SYP*AWNVLPFVCILFYFIEOWF
1		1			1	IVLLEEVFHIPCKLDS
15400	45867	D -	15589	1	2850	IVLLEEVENIFCKLDS
15499		B C	15590	1	2835	
15500	45868					
15501	45869	С	15591	78	491	l

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
15502	45871	A	15592	178	5721	MAVLSLPILAPPTLMPPAHENLY EMQYKATLKEY WORKSPGIPA GANRKKKINGSSPDTATSGGY! SPGDSATGIYGEGHASSATLE LERQPTPSISPNDCSLYPSOS SSNSSSLHAPHSQYRELAVALH SSSAIISQLSENINSLNNENKSAL QLERQVKELKLGELKETVTSDP SNNGWEAVTSLWEGERLPASG AEEPRGSESTAAARPVPVSPAA VRGRLVAGEPGSYQPAETAAK AQLSLMALPGEGY
15504	45872	В	15594	52	855	
15505	45873	c	15595	1	1698	
15506	45874	Ā	15596	302	617	NLAVNPSGPGLFLIGKLLITASIS EHVIGLFRDSTSSWFSLGRVYV SRNLSVSSRFSLVFDDGDVQMG FWCGCPLCLLVFLLT\AGPSAA GLLEFAGGPLQTLFA
15507	45875	A	15597	1	2762	MYGTGYRDVAGKWAIDPDSK REFLDVTQEGIQGCDFSGTCRQ TLSILTQPLRQWGLEGIKKPNS WIISEESVSNGGPPLLIPRQTAS GYDLQQTPTDLQLRVLTVRRK TMKQKGIASTSTKRTSTPKPHL YVTIIKDQSYIKPQRWGKNIAE KLKILKIRVALSLQRNAAPHQQ WNKAGRRMSLMSSQKKASEVI ESQMMEIKGEEKFREKRVKRNE QSLQEIWDYVKRPDLRLIGVPD SDGENGTRLENTLQDI
15508	45876	В	15598	1	1989	
15509	45877	В	15599	1	762	
15510	45878	Α	15600	686	850	GQPERKVGLPTKGSPSD*QRISR QKLYKPKESGGQYSTFLKKRIF NPEFHIQPN
15511	45879	A	15601	2	382	
15512	45880	Α	15602	1	993	
15513	45881	А	15603	3	460	SPCWGQLLQATQP/ATAFAQXD LLVGAVLFGHNGSLSSRNTLAF SMDVGPSQWMWGLLNGCGAS AHRQCTRCTLQVTIALEGPLAP LKLAWHKKIYJFLGEAAEIRRD QLEAERELFHSRAEQGRLFVGR PGQAAHLMLPPGDCVSIWTQH MIT
15514	45882	Α	15604	1	130	
15515	45883	В	15605	1	561	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ļ ·			sequence	1	
15516	45884	Α	15606	94	420	EKAVGRWIFQSKRYRSPCEKPY
l						SKGLQVQRTVPEGPSKAARAA
1				İ		QAPGPNLLPCLPT/PQTPEFIISEP
1						LANMYSCGNQNTLMEELAEQA
		l				QQHDEMLHMHHALKEALSIIS
15517	45885	В	15607	46	747	
15518	45886	Α	15608	186	364	AFAPAPCTLVPGVLRGPPILLRE
		1				TGLAH*CPAVISPPKSLPWLPTA
1		l				YSVESQQVILFGL
15519	45887	Α	15609	155	511	GAAVAGTQNLCLGGGGAVGG
1		1			1	S*RGALNGEGLLHPGYNAPFON
				i		TAATRORGIAQTAPEAAFOLFL
						HQRVRFHC*FLLWREGRGR*AR
1	l	1		1		REPLSRPQPPAGPLGGALPAPPL
		Į.			}	PDASTLVGVG
15520	45888	Α	15610	1352	1684	MSIRSAWCRAEFNSWISLLTFC
13320	13000	l'`	13010	1332	1004	LVDLSLVFDDGDIQMGFWCGC
		ł				PFCLLVFLLT\AGPSAAGLLEFA
l .		1				GGPLQTLFAWASAVVAAEQQI
l .		1				
16621	45000	<u>. </u>	10011	98	367	LVSSKCCCLIVPLEVLSQRSTWP
15521	45889	A	15611			LIGHTION PROPERTY AND A LOCAL
15522	45890	A	15612	426	738	VCVWGDPPGPAGGHGEAAATL
	1	1				WKEKVDLKERVEKLELQFIHLS
ŀ		l				GQTDTIGRKYISQGAVSETQHW
1		l				ERRTSSGWP\GPGGDEGEPAGA
						AGQVLQLVGDHKEGHGKF
15523	45891	В	15613	86	1411	
15524	45892	A	15614	950	1493	GTVFLWRRRGTLIFRVSGFSAL
	1	l				FFPPSLWFYLPFVFDDGDLQME
1	1	l				FWYELSFLLMLLLFLCACWFSF
l		l				*QSGPSAAGLLEFAGGPLQTLF
	1					AWVSAAVAEEQWILVNRKCCC
	1	l				LIVPLEVLSQRSTRPCEVSVHPY
ı		1				VHCCLGIEELNIYCSLHSLGLFE
1		ŀ				AILLGKAFEELGCCDPSLFATV
15525	45893	Α	15615	1	5731	MDSHRPTAWPAAVLAGRSPVA
						AESERYCSRQPPLPRRQALLSV
		1				LVLASDTRLPFPWSSQTFCGILS
		1				KPLVLSGPLOTQGETNNENGLV
		l	i			GGGEVGMOPLVLFERSLFPGRC
1						CVLERRIVPWMRQESYSSSSPI
1	1	1	1			WFVDSDELNLTSVLECLEDDK
1			l	l	1	NNNLLHQQLKWLICGLCRLYN
1			l	I	1	LPKHLDVAMPDQPLPMGQSGP
1	1	1	l	I		SAAGLLEFTGGPLQTLFAWVSA
1	1	l	1	I		
1	1	l	1			ALAAEKQILVNCKCCCLIVPLE
15505	15004	ļ	15616	-	501	VLSQRSTWQYELKEYW
15526	45894	A	15616	2	521	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			ļ	sequence		
15527	45895	A	15617	1	1261	MSEHNITEECOOENOSIEOWRSI
		1				AHSDSIEVSWLHEVIKTYLKAK
						STKLDLMKKLKELSSOPSRKPK
		1		l		PFIPSQEGTGDNDVIYSTERLSK
		l				LLNQKKGSTRLYDKPLSNLLNQ
	l					KKVSTRTEGLTVDSRADMRNV
	l					TTILLSTESSSTDSSSSMDSSSSM
				ŀ		DSSSSIGRVETATCAMIPVQKSK
1						RSGPPQGPSAATGDGAEGRFLP
ł	I					RCSRPWADRLAWSQPRRHLAP
l	ŀ	1				GPSAATGDGAEGRFPPLCSCCG
		1				TDRLAWPQPQGHLALLPRCGE
		l				CGRARELPGDCCLHAEGGCSSV
		l		1		LEVLARAIRQKKEIEEIQIGKQE
		1				VKLSLVADDIIFCLENATDSSRN
1		1				LLELIKEFSKVSRYKINGHKSVA
1		1				LPYINSYQAENHIKNSTPFTIAA
		1				TNNNNKK\KNLGIYLAKESKDS
		1				YNENYKTATERNHRWSQAR
15528	45896	В	15618	60	499	
15529	45897	Α	15619	134	1416	RTSICWTMDSSVQNTVLGPTRS
		l				SVKGVMEWLQYLQMERR/CTL
	İ	l			i	SISMESMPRVAEDAGDVGAAE
	İ	l				PLHLEPPAGPLEVEQQLQAEIK
						HLRKTLESLAGQLQAQVEANE
		l				GSSHLNRQQDERLREWKAELW
		l	İ			EEQGETGRSSVQKDRTTISHAL
		l				SQNHQLEEQLAKERLEATSQQ
						NQQLQSQMSLMALPGEGDELG
	į.					SKEEEEAPQPMPSILEDLKSREA
	i					MVAFLNSALVSAEEEQQAQLH
		l				GQLKEQRVCCQHLAHPVALAQ
						KEPEAAAPTPRTGGDSAVWTV
	ŀ	l				DLQVEKTKFIMFTIKPGEVVNK
	1	1		1		DMAQSIYRSGKNLDKDMYGH
	1		l	l		DLEARRKTNRPLETTFYFLSYE
						LDYSRCFIGCLKCLCLPSPSATI
	1	1		l		GSFLKPSQKQKPVCLLYSLQNH
				l		EPIKPLFFINYPDSDSYRHTQVIR
	1	1	l	l		ILLSGSIHVQVPATKTPL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequenec		
		<u></u>		<u> </u>		
15530	45898	Α	15620	1	1665	MGGHVGGRHRMLTPKLIPETL
						AAGSMKINPEEPNRPPINGAPG
			İ	1		GPALPLAAGDSGSEQAGEPALA
		l				ASELQGWDSTLGSPGSLRGTGI
						VATRSGSRGFRLSPASLGRRSR
				ł		DRKTKELAFAGSGLATRGQRA
i						SSDAAAGPPLSRCSPARTPQLPV
1			ł			RGSLTGPRAPAPPRARVCLLSC
1		1				SCROPFPPLTGGLNVGRPSGPS
1						WKICPRSLPSASAETGIRRRRAA
1		l				AHPPLQQPPLCGESLPSVQCSA
						NADSSIPVPSRWSODACRNTGA
1		l				AGPPGYPACVNNRNRTDKNSQ
1		1				LDETHHMKTKVTDRGGFGAFT
1		l				MLLRVVDTGHTSCSGSVEWDP
						RLYGDQADLSRSGHLDRPGAH
	1					HDYYTAIKKCEEGKSLDTTVK
		l			i	VKAGVNGFGRIEHLVTKPAFNS
		l				GKVDIVTINDPFVDLNNMVYM
		l				FOYDSTHGKFCGTVKAENRKL
	ĺ	1		l		VASGNPITVFQ/ERDPSKIKW\YI
		l				AESTCIFTTMEKAGADLOGGA
	ŀ					KRVIISAPSADAPMFVMDVNHE
		l				KYDNSLKIVSNASCITNCLAPL
						ANVIHNEGIVEGPMTTVHAITA
	İ					TOKIVDGPSGKLA
15531	45899	A	15621	2	1105	GRVGCSTVSRIFSCVASRATSLR
13331	43077	<u>ا</u> م	13021	ľ	1103	TPMGKVKVGVNGFGRIGRLVT
	ľ	1				RAAFNSGKVDIVAINDPFIDLN
			ĺ	l .		YMVYMFQYDSTHGKFHGTVK
	ŀ		ł			AENGKLVINGNPITI\FOER\YPS
		1		İ		
						KIKWGDAGAEYFVESTGVFTT
						MEKAGAHL\QGGAKRVIISAPS
						ADAPMFVMGVNHEKYDNSLKI
				į.		ISN\ASCTTNCLT\PLAKVIHDNF
		l		1		GIVEGLMTTVHAITATQKT\LD
				1		GPSGKLWRDGRGAF\QNI\IPAS
1				l		TGAAKAVGKVIP*AERGKLTG\
	}					MAFRVPTANVS\VVDLTCRLEK
1						PAKYDDIKK\VVKQGVGGPPQG
1						AILGYTEH\QV\VSSDCNSGPPV
1			l			FPPFDAG\AGIALNDHFVKLISW
						YDNEFGYSNRVVDLMAHMAS
15532	45900	Α	15622	121	221	
15533	45901	Α	15623	189	538	

SÉO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
15534	45902	-	15624	3	1015	DRACEDO A DRACO CALIFORNIA
15554	45902	Α	15624	3	1015	DRAGPPGAPRLGQQLLCAIILL
	l .	1				GLAAKKKEINITIIKQLLKKFLN
		1				KKPSYGLERQMETTQNLVDSY
		1				MAIVNKTVWDLMVGVMPKTI
	i i	1				MHVMINNTKEFIFSELLSNLYS
		1				RGDQKTLMEESAEQAQWRDE
	ŀ	1				MLRMYHVLKEALGIIGDINTTT!
	l .	1				ST\PWGPVDNSWLQVQSVLARR
	i i	1				R/SPGLAPTAPSLPTPHRLSLGA
	l .	1				LGTGSVPTLLHHPGSSSVSSLSG
	1	1				LVA/DCFGLPKARGGPGQRLPF
1		1				PSLPGRLFCDLAVAPPAPKAAP
İ		1	l			FICDFVYCCDQASEPVGCQGQT
		1		1		APHRLHGSPLGIPQAPPGSIQA
i		1		1		WLRAAVFWKGALCPRCGRHIQ
		1				EFLLHRCTLVSL
15535	45903	Α	15625	547	1242	PKALIKTSNKSSLSYQLTTHSGT
		1	l	ĺ		MACPWIRHEVLLSRLGFLVLEN
		1				TNMDKRPSKLLRWLERQMETT
		!	i			QNLVDSYMAIVNKTVWDLMV
		1				GVMPKTIMHVMINNTKEFIFSE
	i	1				LLSNLYSRGDQKTLMEESAEQ
		1				AQWRDEMLRMYHVLKEALGII
	l .	ĺ				GNINTTTIST\PWGPVDNSCLQV
		1				QSVLAGCSFIILVQAVFLLYQA
	i i	1				WWLLFWAPQGERWPWTSGLE
		1				DTVTREEGSPKGLSISL
15536	45904	Α	15626	108	308	AQKHHLT*LSFLPRH*ILT*NLT
l	1	į.		i		KIPTSKIKKTPRSRSPRRSTSCFL
		1			İ	DNLAPSCGCSGKGSSWRIC
15537	45905	Α	15627	687	1910	KDHQLRKKLRRRGKQGAFTAA
l		1				LPPGSPRPPLRGGRSGGVGLGV
1	į.	1				RGQRTPQAPQAVTPSEGPEPLW
	1	i i	i			GPVLGGGGEHRRGTGASAAAG
	I	İ				ASMRRLGAAGGGAGAGPORG
	1					VGARVVVSEEANSPGAAITSNR
						ETQDANEGLSAPRPECLLHPTC
						RLLYVPLIVRDWVCHTRSVVLL
						QVVNVTGIAVGTGLASAYDTL
						MSOSFGVKNLKRVGIILORGVL
						ILMLCCFPCWADLVNTERILLL
						LKQDPKSPGWTRECFQEWGSYI
						HLVIPSMFMVCTEQWTFEIGNF
l		1		I		LAGLIDVMELGTQGIICELASV
l	1	1		I		AYMNYFRLNPQPPISDFTINLAF
						LTLTERKILGYIQLRKGPDIVGP
				İ		YGLLQPFTDAVK\FTKEPLWPS
	1	1	1	I		TSTITLYIIAPTLALSIIVLLALPT
1				l		RMLAATPRPRGT
15538	45906	A	15628	1	1173	MADE ATTEM NOT
15539	45907	A	15629	3	1447	
13339	1.3707	1′	1.5025	Ľ		

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15540	45908	A	15630	1	215	MDPKQEEIPDSPEKEFRRYRIAE WIRIHQPIICCLQETHLTHKDSH KLKV/KGWNKAFHANGHQKSA ARETK
15541	45909	Α	15631	1	1524	
15542	45910	A	15632	1	575	MALSERSOPOHYSHIDRIYVK HPEYANLYROKVDYTLPRVSIS HIVGQAGVQIGNARWELYCLE LAIQPDGQMPSDKTIGGGDDSF NMFFSEIGAGKHVPRAVFVDLE PTVVDRCLGHWMAAFLGEEV HTGTYRQLFHPEQLISGKEDAA NNYSKGHYTIGK/EDLVLDRIH NLYSKGHYTIGK/EDLVLDRIH
15543	45911	A	15633	208	465	KEETSQCIALITETTII WESQITT
15544	45912	Α	15634	107	827	EARRPCTLHLRSALHPATMCEL YSKRDTLGLRKKHIGPSCKVFF ASDPIKIVRAGRQYMFDENGEQ YLDCINNVAHVGHCHPEVVKA ALKQMELLNTNSKFLHDNIVSKFLHDNIV AKRLSATLPGETLCLLFYKFRG YWFSSTNIPSSRPAAPASPPAAE RAGDFRAGRESVPGSSRVSGS PRGPCSKLSLSTFCTRYCPLSAL TSGGCCPYMCFFLSPR VSRLLY SSHMVAGCRALRCKNSH
15545	45913	A	15635	3	1218	GGLWEKMAAAA\GRVVRYLS MSRSAITALTSVCHIGPPCRQL HHALMPHGKGGRSSVSGIVAT VFGATGFLGRYVVNHLCTAW GQQVIIIPIYRV*IKY*QSMHLSFP WG*PGGQLSVFLGNWGTARD* RFLFRRVSYNTGNVGSSILFGTR LGETKNFLOFWRDVFVKIPQAIA VQLSKGSWEFEKFISCFHIWNG IFKALS*01*GNKA VGEKVVGD AFPÆAIIVKPSDIFGREDRFLNS PASMHRFGPIPLGISLGWKTV KQPVYVVDVSKGIVNAVKDPD ANGKSFAFVGPSRYLLFHLVK VIFAVAHRLFLPFPLPLFAYRW VARVFEISPFEPWITRDKVERM HITDMKLPHLPGLEDLGIQATP LELKAIEVLRERRITYRWLSAEI EDVKPAKTVNI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15546	45914	Α	15636	2	799	GLWEKMAAAAQSRVVRV\LS\ MSRSAITAIATSVCHGPPCRQL
		1				HHALMPHG\KGGRSS\VSGIVAT
1		1				GFGA\TGFLRRYVVNH\LGRMG
						SQVIIPYRCDKYDIMHL\RPMGE
						LGQLLFLEWDARDKDSIRRVV
						QHSNVVINL\IGTRP/WKSKNSD
						YEDVFVKIPQAIAQLSK\EAG\V
						EKFIHV\SHLNANIKSSSRYLRN
						KNRKREKEEMKKDLLSKKEPE
		1				PEDLENSQLIHIAKNEKACFEEN
ł						TKSVAEQPFDEEIVDATHQLTS
						YLGRSQE
15547	45915	Α	15637	560	1394	RKSSSVHNAASGLLGSRAPGAH
ļ						RHLGRCRCRHRLVSPSPAMAE
						DLSAATSYTEDDFYCPVCQEVL
						KTPVRTTACQHVFCRKCFLTA
						MRESGAHCPLCRGNVTRRERA
						CPERALDLENIMR\KFSGSCRCC
						AK\QIKFYRMRHHYK\SCKKYQ
						DEYGVSSIIPNFQISQDSVGNRIE
1						NNTGIKQIPGAPAFGDGRVGCR
1						QRNAKTELLAAAGESPCTMSL
ŀ						GSGPNLGPSPRSKSSSVTSPQPC
1						FTEHPRERFVSFPVHYVLEGSR MSFLLWSLKTDRIFGH
15548	45916	A	15638	1	1026	MKVVCKEVKFEKOCPEAVLLH
13346	43910	^	13036	1	1020	WEQTDDDENKRCDLREPRKPE
ŀ	į					THTKGPGLLVPTTOPGRNEGOS
	į					GADGKHARCSKTSCCLSGKEPS
	ĺ					FIITEHCPLVPVQNTTIKPDSGIR
						ILQTLDSRETFTMSTALDVLQM
				ŀ		KEEDVLKFLAAGTHLGGTDLD
						FRMEQHICKRKSGGIYIINLKRS
						WEKLLLVAPAIVAIENPADVSV
1						ISSSSTGORAMLKCAAATGATP
l						IAGHFTPGTFTNQIQATFPNPRV
				1		LVVTDCRTGHQPLAEASYV/NN
			1			LPTIALCHTDSPLHHVDIAIPCN
l				1		R/KAHSVDLMWMLAREILPM\R
				l		GTIARKHWWE\DPEEIEKEEQA
			l	l		AADKALTKEEFQGE
15549	45917	Α	15639	20	441	TAEIAIPCNNKGAHSVG*MWR
			1	l		MLAREDLRMRGTISGEHPWEV
1		ļ.	1			MPDLYFYRDTEEIEKEEQAAAD
						KAVTKEEFQGEWTAPAPEFTAT
	1		1			QPEAADWSEGVQVPSVPIQQFP
l			1	1		TEDWSAQPATEDWSAAPTAQA
		L_				TEWVGATTDWS
15550	45918	В	15640	88	531	
15551	45919	A	15641	1	1302	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15552	45920	A	15642	150	957	IKAGASCCGSANPYVSVCKSCV LLAMAQLQTREYTDNKKYAVD DVPLSIPAASEIADLSNIINKLLK DKNEFHKHVEFDELLKGQFLRM PLORHMEMENISSEEVVEIE EKYTAPQARSNAWFHDDWISSI KRGRGRILTGSYDKTSRIWSLE GKSIMTIVGHTDVVKDVAWVK KDSLSCLLLSASMDQTILLWEW NVERNKVKALHCCKGHAGV NSIAVDGSGTKYFGLRIEPKPA VSLVEEINSLSYRWTADNYKV GSGGKEEED
15553	45921 45922	A	15643	335	537	VFCCPHCSLTFSKSFSSFSRLVR HQQTHWKQKSYLCPICDLSFGE KEGLMD/HLEGL*RQOFGQSSH HKCRVILGQWLGFSHDVPTMA GEEWKHGGDQSPPRIHTPRRRG LREKACKGDKTKEA VSILKHK VYFKQVTLDSSCRL*ERQVKST
15555	45923	A	15645	153	828	SSWNILSSTWRLPFCPTSVVLRF LTWICLRTISTSSISWFPADSS

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SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
10000	140004	-	15646	I	2002	MLESAGNQDIEDVDIVLRQIQV
15556	45924	Α	15646	1	2002	
1		i				KNRSTTEVGQNGNLQVEVSYT
1						QAHIAPRPLASQGTWPHFVSSA
	l	1	i			PQPCFQVQNHSKYQNLYYSGFI
						RETEPISKGDVGEWLSAGKGES
ļ						SAMFASEQEISKDEQGTPVLGS
						FYWEVDSPRKESSQAWAPGQE
				l		WIKLERDTTEEKMFEQLKPIEP
l					İ	VQKTLPWVGEVAATLQEAMK
				I		RDCWREARVKKKPVTFEDVAV
		ı		1		NFTQEEWDCLDASQRVLYQDV
		1		İ		MSETFKNLTSVEGKKKELREQ
		l				HPSLRDEGTSDDKVFLACRGA
1		ı				GQCPLSAPAGTMDRTRVLQAS
1		l		1		QAGPP/FFFATTVGKCFSRRSYL
		1		1		YSHQFVHNPKLTNSCSQCGKLF
		l		1		RSPKSLSYHRRMHLGERPFCCT
		1		1		LCDKTYCDASGLSRHRRVHLG
		1				YRPHSCSVCGKSFRDQSELKRH
		1		1		QKIHQNQEPVDGNQECTLRIPG
		1	İ			TQAEFQTPIARSQRSIQGLLDVN
		ŀ				HAPVARSQEPIFRTEGPMAQNQ
	l					ASVLKNQAPVTRTQAPITGTLC
		l				QDARSNSHPVKPSRLNVFCCPH
				1		CSLTFSKKSYLSRHQKAHLTEP
1						PNYCFHCSKSFSSFSRLVRHOO
	İ	l		1		THWKQKSYLCPICDLSFGEKEG
1		l				LMDHWRGYKGKDLCQSSHHK
						CRVILGOWLGFSHDVPTMAGE
		l				EWKHGGDQSPPRIHTPRRRGLR
i		l				EKACKGDKTKEAVSILKHK
15557	45925	Α	15647	1	1048	MPSLAELFEGSSKAFGAAASQL
						NFPVASGEMLKFKYG\LRNPLH
1		1		I		AVAAEPIASRTSRLNLFFQGKPP
		1		l		FMTQQQMSPLSREGILDALFVL
		l		l		FEECSOPALMKIKHVSNFVRKY
		l				SDTIAELQELQPSAKDFEVRSLV
		l				GCGHFAEVQVVREKATGDIYA
		l				MKVMKKKALLAQEQVSFFEEE
		l				RNILSRSTSPWIPOLOYAFODK
		l				NHLYLVMEYQPGGDLLSLLNR
		1				YEDQLDENLIQFYLAELILAVH
1		1				SVHLMGYVHRDIKPENILVDRT
		1		1		GHIKLVDFGSAAKMNSNKMVN
		1				AKLPIGTPDYMAPEVLTVMNG
1		1	1	1		DGKGTYGLDCDWWSVGVIAY
						EMIYGRSPFAEGTSARTFNNIM
	L	_		L	L	PINIT CYCLI VEG LOVE I LIMINIM

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	ļ	<u>L</u>				
15558	45926	A	15648	101	220	KRQQAFGLLDLHQWFARGSQG
						FRLQTEGCIIGFPTFEV*RHSGFL
		l				SCQSSCTGSFSSVWADVPLMFE
		1				GAAQPV*PVPSRGSADFRPLSIC
15550		I	15640	***	000	*RSSRILLVKRCENSGVR
15559	45927	A	15649	597	808	ITMKSQRQIIKKAVTTKEMKSV
		l				CRRD/TPAFMFIAALFKIAKIRIQ
		l			1	RLCLSTDEWIKKMWYIHTMEY
15550	40000	_	15650	161	0156	YSTLK
15560	45928	В	15650	161	3156	
15561	45929	A	15651	14	270	
15562	45930	В	15652	787	1821 2412	MANAGORA ATLANTION VID
15563	45931	A	15653	1	2412	MAKSKGCSAATLASTHQILVTP
		l				WSCDNQKCLQTLPNVPGGTKL
		l	1			LLCEDHWFRNDSCNQAIPTIAT
į .		l	l			LLSFPSLQASLWTTHFLSPLNH
		1				HPVNPDSHLEQPPGQSPRLNFE
		1				AGDPERAPHALGCKRSRTRATS
		1				DRQPMASRDPGWGKASVGKA
		1				CALGSAVLPAGNDCTTDNLGL
l		1				GSAGFEDLRLRLGQGPPRRQGA
		1				LQAAPRKASSCAAGQRLAPDSS FSRKRRPSRQRLGCETAAVADT
l		l	İ			AEAKSSEAPAFGWSTMDEAGS
		1				
		1				SASGGGFRRALDEPPNSRIFLVI
		l				SKYTPESVLRERFSPFGDIQDIW VVRDKHTKESKGIAFVKFARSS
l		ı				QACRAMEEMHGQCLGPNDTKP IKVRVPGSGCPRGKEWASWTSL
		l				HLLLCRGEGGVEHPFPAADQT
		1				APGLGTGAALRRGSPPTKLRRV
		1	1		i e	AGKGTGCGVFAPRERTROPRPP
		l		l		GDRTAVAGLP\SPSPSLIHQLLSS
1		1	l	l		
		l				LPNSS*PTLSTFLTFLSQMPRFP
		1	l	l		YIKNGKTL/PFPQVFIAQSRSSGS
		1		I		HRDVEDEELTRIFVMIPKSYTEE
		1		I	1	DLREKFKVYGDIEYCSIIKNKVT
	ŀ	l				GESKGLGYVRYLKPSQAAQAIE
l		l				NCDRSFRAILAEPKNKASESSE
1	l	I	1			QDYYSNMRQEALGHEPRVNMF
1		l				PFGHGVVQYFNVASAIYAKYK
		l				LHGFQYPPGNRIGVSFIDDGSN
1	1	I				ATEKMATQMVAAQLASMVWN
		<u> </u>				NPSQQQFMQFGGSSGSQLPQIQ

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	scquence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15564	45932	Α	15654	375	2059	DSSVWOTPOKORAVRPLHSGG
						APWTKLAALRATRGFRPGVDS
						LDEPPNSRIFLVISKYTPESVLRE
l						RFSPFGDIQDIWVVRDKHTKES
						KGIAFVKFARSSQACRAMEEM
						HGQCLGPNDTKPIKVRVPGSGC
ĺ						PRGKEWASWTSLHLLLCRGEG
						GVEHPFPAADQTAPGLGTGAA
				l		LRRGSPPTKLRRVAGKGTGCG
						VFAPRERTRQPRPPGDRTAVAG
						LPLSLALTNPPALEFTAKLVLA
			ŀ			YFEYFSDLPLSDAQISIYKKRKN
						STFPQVFIAQSRSSGSHRDVEDE
						ELTRIFVMIPKSYTEEDLREKFK
				1		VYGDIEYCSIIKNKVTGESKGL
		İ				GYVRYLKPSQAAQAIENCDRSF
l						RAILAEPKNKASESSEQDYYSN
l						MRQEALGHEPRVNMFPFGHGV
						VQYFNVASAIYAKYKLHGFQY
		1		l		PPGNRIGVSFIDDGSNATDLLIK
						MATQMVAAQ/RLHQWCGITQV
						SNNL/WQFGGSSGSQLPQIQTD
						VVLPSCKKKAPAETPVKERLFI
						VFNPHPLPLDVLEDIFCRFGNLI
						EVYLVSGKNVGYAKYADRISA
		L.				NDAIATLHWKDSEWGET
15565 15566	45933 45934	A	15655	257	412 866	IVA A SPECICLE PROCESS OF THE PARTY OF THE P
12200	45934	Α	12020	3	800	HAASPCCCYEEKQSPLQKHLRV
						CSLRTWSLAARVAEGTAETVD
						PSAAPKTSVWSSRVCWGRLLA TISPSVNETDTCGMEDRRLVVC
						RPWRAOCHMKAWGWRHPEEG
						GTHSGAPSAVLQALAVAIQLGG
			ĺ			HLADPLLOVDPLSSCGAVSLDI
		l		•		LIYLVFYYRTASVPETYIVKTLF
l		1		l		KKLESQSLIQSNVLTRSNSMKA
				l		ERGEEAAKEKSEASRGWFMRF
		1	İ	l		KERSCLHNIKMOGEAASADVE
l		l		l		AAASYPKDLAKITDEHGYTKQ
				1		O\FDVYKTAFCWRVMLKKYVK
	l			L		LLFVLCPLP

SEQ ID	SEQ ID NO:	IM.	SEQ ID NO:	Nestantida	Nucleatide location - f1t	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
	1	ı		sequence	' '	
		_	<u> </u>			
15567	45935	Α	15657	333	1034	RLSTTVMRGAFVF\MHKMPRL
		1				WLDYCQF\LMDQGRVTHTRRT
		İ				FDRALRALPITQHSRIWPLYLRF
		1				LRSHPLPETAVRGYRRFLKLSP
						ESAEEYIEYLKSSDRLDEAAQR
						LATVVNDERFVSKAGKSNYQL
						WHELCDLISQNPDKVQSLNVD
						AIIRGGLTRFTDQLGKLWCSLA
1						DYYIRSGHFEKIRKAPVNGGPT
l	l					WMVGQNSVPRFQTRRLTPLKR
				ŀ		KGPLVTDFREKQPTPLAQE
15568	45936	Α	15658	172	346	ARPCRWCPGS*TSASAPGPGLR
						PCPP\GGDRTPASPPLHNQAAA
	1					APPGPGTWRGAAAS
15569	45937	A	15659	293	482	
15570	45938	Α	15660	468	791	LAKCGCLEARGRARQTSGRPRP
1						TDGGP*SPCWETGKGHPVLGR
1		1				RGS/PGKQGL*SLPQSCQGQWR
1		1				CAASRLLSSKNSWACMGSREA
		l				VSQETGTSSSSLGLMRPSDKKP
15571	45939	Α	15661	366	609	
15572	45940	В	15662	321	739	
15573	45941	Α	15663	1	1815	
15574	45942	Α	15664	1	1911	
15575	45943	Α	15665	437	1125	PVSCSGHSRWAAAAISIALSMIE
ł	i	}				TVNMVFTFYSRHHGTISTRVL/I
1		-				GKRPERODGODTVNLGTLEET
ŀ	ł	1		1		ASPSPVVTTSPPOPMLPSAFPPL
1		1				SEEINPALPEATVMASPEAVAR
	i		-			ODNVDSAQEPPPTPLCASRPTT
	l .			1	i	ROKSWWGARGEVESVTHEEM
				İ		HYTGKELLEFSNLYKEKSGEQA
1						WEWILRVWDNGGRNIOLDHPE
1	i i	l				FVDLGPLSRNSAFNVAVRGVK
ŀ	į.					KGSKSLFAWLGEI
15576	45944	c	15666	49	186	KOSKSEI AWEGEI
15577	45945	Ā	15667	80	877	GAERLPEGPGKRMVFRRFVE\V
13377	43943	l^	13007	00	0''	GRVAYVSFGPHAGKI/VSAMVD
		1	l			VF\DONKGFWSNGPCTQVR\RQ
		1				A\MPFK\CMOLT\DFILKFPHSAH
1						
		l		1		Q\KYVRQSLGRREDINTKLGQP
	1	l				TRWGQGRFESQEEKEKPRLTDF
	1	1		i		*SVF*S**RAKEN*RDRINPRNE
						V*EAFKRAAFPGKASSQKKAPG
	1			l		Y*GVLLAAAAAAAAAAAAA
	1	1				AAAAAAK\VPAKKIT\AASKK
1	1	1		l		APAQKVPAQK\ATGQK\AAPAP
	1	1		l		KAQNGSKSSKPQKAPAPKASG
						KESISGNHKK
15578	45946	Α	15668	253	1218	
15579	45947	Α	15669	I	1819	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15580	45948	Α	15670	1	2422	MCPILREHRGSTEGKRRQKRLP
		l				LELTLKDEEVLAHRQEIEVAKA
		l				SPSKGNIIIHFGYKGVEYHDQQ
		1				DVTSNFLGAMWLISITFLSIGYO
						DMVPNTYCGKGVCLLTGIMGA
						GCTALVVAVVARKLELTKAEK
						HVHNFMMDTQLTKRHCPWAA
				1		TIQPKLSFLISLPGCDYRGESAR
		İ				ARPLSHPPLQLGLGKDARYLRI
1				1		CDRSLLCVCAAAGNAGLRPRD
				1		SPHTLLHWRSRSPYFFSPKPVPP
		i		1		QLLGGQPAAHSRRDDLGEAGV
				I		GRGGCCFAALRGVQSEKWSGF
		l				HTVPEPAPGQGDVMPIVLVRPT
						NRTRRLDSTGAGMGPSSHQQQ
		Į.				ESPLPTITHCAGCTTAWSPCSFN
						SPDMETPLQFQRGFFPEQPPPPP
		1			İ	RSSHLHCQQQQQSQDKPCPPFA
		l			Ì	PLPHPHHHPHLAHQQPASGGSS
		l				PCLRCNSCASSGAPAAGAGDNL
		l		İ	ł	SLLLRTSSPGGAFRTRTSSPLSG
		ı		¥.		SSCCCCCCSSRRGSQLNVSELTF
		l	l			SSHASALRQQYAQQSAQQSAS
1		1				ASQYHQCHSLQPAASPTGSLGS
ŀ		l		l		LGSGPPLSHHHHHPHPAHHQH
[Į				HQPQARRESNPFTEIAMSSCRY
		1				NGGVMRPLSNLSASRRNLHEM
		1	l			DSEAQPLQPPASVGGGGGASSP
		l				SAAAAAAAVSSSAPEIVVSKP
		ı				EHNNSNNLALYGTGGGGSTGG
		1				GGGGGGSGHGSSSGTKSSKKK
		<u> </u>				NQNIGYKLGHRRALFEKRKRLS
15581	45949	A	15671	1	246	SEPPHRGPTGALPSGAVRRGPL
		l	1			SSRPQIGRSTDSLHCVPGKATD
						AQRQPVKAAGSGAVPCKATG
15582	45950	В	15672	43	741	WSCPKL*EPISCINVTIC
15583	45951	A	15673	687	1012	FHTSRPFWNISELSRFCIQKYKK
15565	43731	<u>۱</u> ^	15075	087	1012	PWHSVSDFYRYOFFICOGTACC
				l		PASLRM*HCPN*ILQGTSSQNCC
				l		SCSGKSCSPMQQRHRSTVCSGS
				l		GTFCVGSCESFEGTYFSR
	L			L	L	OTTO VOSCESTEDITESK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15584	45952	A	15674	1	1640	MSCDTNGEVEGGRVDSKTRGR
						QEVERNTETHLGSGLNCDIPLIF
	1	ļ				LTYEWRGILEYKIFSVKTGEVL
						GKPGQLGHPTGQTTFFETGTQV
	1					TSMQEMGSYSFGQPHPVALQG
	ļ.					TAPLPAAFTGWRLASVAFPGTQ
	ŀ				i	CKLSVDLPIWGLEDSGPLLTAP
	i	l				LGSAPVGPLCGGSDPTFPFCTA
		1				LTGPSAASLLEFARGPLQTLFA
						WVSAEVSAEQGFFVNHECCCLI
	1					VPLEVLSQRSTRPCEVSVCPYW
						GVPPRDRSNQCPSGFTLDSVGP
	I	1				FCADEDECAAGNPCSHSCHNA
		ŀ				MGTYYCSCPKGLTIAADGRTC
	1			l		QDIDECALGRHTCHAGQDCDN
						TIGSYRCVVRCGSGFRRTSDGL
		l				SCQDINECQESSPCHQRCFNAIG
		1				SFHCGCEPGYQLKGRKCMDVN
	i i					ECRONVCRPDQHCKNTRGGYK
	1	1	l			CIDLCPNGMTKAENGTCIGECL
		l				AVSVTEIS/CAVNKSNIDECKDG
		1		ŀ		THQCRYNQICENTRGSYRCVCP
	1	l				RGYRSQGVGRPCMDIDECENT
		1				DACQHECKNTFGSYQCICPPGY
		1				QLTHNGKTCQGKLPFKQIESGG
15585	45953	Α	15675	1	669	MGFCHDGRSIAFRGERGNDESA
		1				IEMIKVSHLKQYLAVVFRDKPL
			1			ELWDVRTCTLLREMSKNFPTIT
		1	ļ			ALILTGRCASLKWRSKPKKEEF
		l				MGYRQWNPTQESSKGDFEIQL
	i	1				EGGPSMATVYRTEMQPVQIGIE
		1				LLPSPLSFHPSPNSGIPNDPSWL
				l		LSTATLHPPTGVERSSSPATEQR
		1				WTENDFDELREEGFRRSNYSEL
				ļ		REDTQTKGKEVENFEK\NLEECI
15586	45954	A	15676	1	323	MAHESAEDLFHFNVGGWHFSV
1		1	l	I		PRSKLSQFPDSLLWKEASALTS\
		1	1	1		SESQRLFIDRDGSTFRHVHYYL
						YTSKLSFSSCAELNLLYEQALG
						LQLMPLLQVRCSCFLGVVDQ
15587	45955	В	15677	1	1443	

SEO ID	SEO ID NO:	Met	SEO ID NO.	Nucleotide	Nucleatide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codun, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15588	45956	Α	15678	1	708	EVTASPOGVSAVSKGDOKWI
				1		ALGKGLTFIKAVGSIORNRWRG
						WOLKALYOCTTIAGATGVFLR
						VYRALDVLOMEEEDVLKFLAA
						GIHLGGTNLDFQMKQYIYKKK
		l				CDGVYIINLKRTWEKLLLAACA
						IVAIENPADVRVISSRNTGORAL
		1				LKFAAATGATPIAGRFTPGIFTN
		1		l		OIOATFOEPOHLFFTDSRADH/S
		l				PLTEASYVNLHTIAPRNTDSPLC
		l				YVDIAIPCKKGAHSVGL
15589	45957	Ā	15679	78	757	REIFINVRSLLMACFVKEEDVL
1						KFLAAGTHLGGTNLDFQMEQY
			-			IYKRKSDGIYIINLKRTWEKLLL
						AARAIVAIENPADVSVISSRNTG
				1		QRAVLKFAAATGATPIAGRFTP
				Į.		GTFTNQIQAAFRE\PRLLVVTDP
	-	1				RADHQPLTEASYVNLPTIALCN
						TDSPLRYVDIAIPCNNKGAHSV
						GLMWWMLAREVLRMRGTISR
						EHPWEVMPDLYFYRDPEEIEKE
		ĺ				EQAAAEE
15590	45958	В	15680	1	1767	
15591	45959	Α	15681	98	412	
15592	45960	A	15682	521	600	PSPGES/PPLQPDPPVPPWPPPEE EPRA
15593	45961	Α	15683	117	285	
15594	45962	С	15684	12	468	
15595	45963	В	15685	1	840	
15596	45964	Λ	15686	1	1173	
15597	45965	Α	15687	1	2931	MDLFMNVAMLRAARYLWSEA
	ļ	1				VSGFGAQDPKSLALRTHCQTSG
						WSLTEQDPYNNVIRTTIEALAA
		1		1		TLGGTQSLHTNVFDEALGLPTD
						FSARIARNTQIIIQEESELCRTVD
						PLAGSYYIESLTDQIVKQARAII
						QQIDEAGGMAKAIEAGLPKRMI
						EEASAREQSLIDQGKRVIVGVN
				1		KYKLDHEDETDVLEIDNVMVR
				1		NEQIASLERIRATRODAASDTR
		ŀ		•		EIVDLTNGNIKTFNGSIIIYARFD TGRHLFSGGNT
15500	45066	С	15688	11	1779	TGRHLFSGGNT
15598	45966 45967	В	15688	22	1531	
15600	45968	A	15690	98	412	
15601	45969	Ā	15691	521	600	PSPGES/PPLQPDPPVPPWPPPEE
						EPRA EPRA
15602	45970	С	15692	175	454	
15603	45971	Α	15693	332	494	RKEMASGFSKGPTLGCC\RRAL
l	l	1				PDGDTQLQLLLRGNHDSVLGA
		Ь.	L		L	KRRDKGEAGPD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15604	45972	A	15694	2	1276	HTLHSLVTILGTWALHSGPALL
						LPRPGSGLDFLLSPVLPSPOPPG
		1				LPTPTPFTNAVOLLLTLKLVSLA
						SEVQDLHLAQRKEMASGFSKG
						PTLGLL/RRRALPDGDTQLQLLL
					1	RGNHDRPVLPLPHLPGLAGAAL
	l .	1				PRGSA\SLRPLLRRAWPAPLFGL
						LFLLSSHLFPLEAVREDAFYARP
		1				LPARLFYMIPVFFAFRMRFYVA
						WIAAECGCIAAGFGAYPVAAK
ľ	1					ARA\GGGPTLOCPPPSSPEKAAS
						LEYDYETIRNIDCYSTDFCVRV
ŀ			i			RDGMRYWNMTVQWWLAQYI
	i .					YKSAPARSYVLRTAWTMLLSA
	į.					YWHGLHPGYYLSFLTIPLCLAA
						EGRLESALRGRLSPGGQKAWD
	i		i			WVHWFLKMRAYDYMCMGFV
	i		İ			LLSLADTLRYWASIYFCIHFLAL
						AALGLGLALGGGSPSRRKAAS
		1				QPTSLAPEKLREE
15605	45973	Α	15695	134	601	RDGEQPLWAWGSPCSPVAPTL
	1					CILWAPSSGPGPSFRPSPAPATP
l			l			WLWPGLSPISCDSEPSASWACP
l						LPCPSPKPDRWLLTLKLVSLAS
1		ŀ				EVQDLHLAQRKEMASGFSKGP
1						TLGLL/RRRALPDGDTQLQLLL
	1					RGNHDSVLGAKRRDKGGQTTII
						FVQV
15606	45974	В	15696	76	504	
15607	45975	Α	15697	11	706	HITRGRSCTRWTSSAMAKGAPR
1						VILLCLFGAALCLTGSRALQCY
	1					SFEHTYFGPFDLRAMKLPSISCP
						HERFEAILSLDTGYRAPVTLAR
						KGCWTGPPAGQDAIEPGRAAA
						RLLG\VRGCTTDKCNAHLMTH
l	1	1		i		DALPNLSQAPDPPTLSGAECYA
		1		l		CIGVHQDDCAIGRSRRVQCHQD
	1	l				QTACFQGNGRMTVGKGLEGW
		1		l		KGLGKGYGGRAAMRVVIPDRP
l	1	ı	l	l		NRETWPCSEVGGMTLV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15608	45976	IA	15698	2	1642	DEGRAKGHHTWOOTRENESOA
13008	43970	^	13096	1	1042	KGETPYKTIRFRETYYHKNSMO
						ETTPMIQLSPTGSLPQHVGIMG
						ATIODKIWVRTGLPGFPGTPGL
						PVSSGFPGSPGFPGLQGPPVSYG
		1				PTGIPGPIGPPGPPGLMVSSGPP
						GLPGPKVILGLNFQGPKGEKVS
						KGLQGPPGPPGQISEQKRPIDVE
		1			•	FOKGDOVSNGDRGPPGPPGIRG
			}			PPVSTGGEKGEKGEOGEPGKR
		ļ.				
		ļ	1			VSEGKDGENGQPGIPVSSGDPG YPGEPGRDG/EKGKNVMGPPGP
1		1	1			PGFPGERGOKGDEGPPGISIPGP
į.		ŀ				
			1			PGLDGQPGAPGLPGPPGPAGPH
			1		-	IPPSKLGDTCFNCIGTGISGPPGQ PGLPGLPGPPGKL\GAPGAPGFP
			İ			GSKGEPGDILTFPGMKGDKGEL
					ŀ	GSPGAPGLPGLPGTPGQDGLPG
						LPGPKGEPVSCTFKGERGPPGN PGLPGLPGNIGPMGPPGFGPPGP
	ĺ					
	l					VGEKGIQGVAGNPGQPGIPGKF
			1			GDPGQTITQPGKPGLPGNPGRD
				İ		GDVGLPGM*GLPGQPGLPGIPG SKGEPGIPGIGLPGPPGPKGMLL
15609	45977	A	15699	246	433	GLFPLEPGANGI*GC/SGRIRAO
1300)	43311	1	13077	1-70	1-33	RWLPGRPKFTGESFIR*PPRVAK
	ì	1		l		ESGQLIWFGCVPTQISS
15610	45978	A	15700	179	5262	RSCGSRRRMKLRGVSLAAGLFL
13010	43576	,,	13700	l'''	3202	LALSLWGQPAEAAACYGCSPG
				į .		SKCDCSGIKGEKGERGFPGLEG
i						HPGLPGFPGPEGPPGPRGQKGD
		1			i	DGIPGPPGPKGIRGPPGLPGFPG
		1				TPGLPGMPGHDGAPGPOGIPGC
						NGTKGERGFPGSPGFPGLQGPP
					· .	GPPGIPGMKGEPGSIIMSSLPGP
			l			KGNPGYPGPPGIQGLPGPTGIPG
			ŀ			PIGPPGPPGLMGPPGPPGLPGPK
						GNMGLNFQGPKGEKGEQGLQG
						PPGPPGQISEQ
15611	45979	A	15701	155	623	FNYLPPGPSHN/DVGIMGATIOD
15011			10,100			KIWGPPGLPGFPGTPGLPGERG
1				ļ		FPGSPGFPGLQGPPGLPGPTGIP
1		1	1			GPIGPPGPPGLMGPPGPPGLPGP
	1	1	1			KGNMGLNFQGPKGEKGEQGLQ
	1	1	1	1		GPPGPPGQISEQKRPIDVEFQKG
1		I	1			DQVSNGDRGPPGPPGIRGPPVS
15612	45980	A	15702	581	687	KKGAGQVV/DDVQLAQRLAQA
13012	1.5560	ľ.	13,02	I		NEIVLSWNCWMLCKQY
15613	45981	A	15703	7	123	
.5015	.5501	۲.	1.5,05	<u> </u>		

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15614	45982	A	15704		374	RFRKAQCTIVERLTNSMMMHG RNNGKKLMTVRIVKHAFEIIHL LTGENPLQVLVNAIIN/DVSPLR RVNHAIWLLCTGAREAAFRNIK TIAECLADELINAAKGSSNSYAI KNKDELERVAKSNR
15615	45983	A	15705	2	892	QAFLEITPDOLSNYKVFAKKTA LWONDDVHSQHKGEVRNAAG EFKYRNSVTTHSEGKWQSEKG KDLGVGYWSMACYKLGHIAG NEQQKHICGGIAKKWKQPKCP PTGKWPAVVKFATESEGGGAK KEEKAHLPRVVEKELHESSRII RHSIESVLLQAFTQKDYTAVKE KYAKYLPIISAERYAAKFEKA QCPIVECLTYSMMTHGRNNRK KLMIMRIVKHAFEIIQLLTGDNP LQVLVNAINSGPREDSTLTGRA RIVRRQAVAVSPLSKVNQAIWP QCTGALEMAFRINKTIA
15616	45984	Α	15706	1	536	
15617	45985	Α	15707	8	437	
15618	45986	В	15708	255	1058	
15619	45987	A	15709	609	1489	LPLSLPRAARPAVAPHLPPREPGPALPLPPPGPEPPSLPGSAGE NS*RTGTAAAGAGPFAPASMSL RVQTPCWPQRGAREPPGSCAS WLFCAVWQMYRAHSGCPLAPP PAAAAWVAAQLQP*DGHEGPG G*ET/S*VLRLRPTPTPPLPTWCC TVRPRRPVLPFCLIMRRRTAGPP CAPIAAAPTTPSSLRRGKVPTGR PPUL*G*PAVMLTMAVTITRTV TTSCNAAAVMPKTGTAHRAP LTPGTV*RDLCLVKPTAQMPPS APPWTAPPRAVAPVIVLPAIS NGAFA
15620	45988	Α	15710	173	530	GRERGDKGGAVQTLQDLVNV HLGLTYPPRPCGHSPPSWASPQ APLQPPLRGEPGHGWGS/PSGL GQGKSEGLGPAHLPRAARPAV APHLPPREPGPALPLPPPQPPRPP SLPGSAQGCNS
15621	45989	Α	15711	1	227	
15622	45990	A	15712	238	498	
15623	45991	В	15713	1	1017	
15624	45992	В	15714	590	649	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540.217	codon for peptide sequence	of peptide sequence	deletion, *possible nucleotide insertion)
15625	45993	Λ	15715	3	843	AVFLSTSNEAVYKIFDDSPCPFT
				l		AKTCNPETFPSSNESRQCPNAR
		l				COFAFYGGESGYHRALLGLOIF
1		l		1		NAFMFFWLANFVLALGOVTLA
				1		GAFASYYWALRKPDDLPAFPLF
						SAFGRALRYHTGSLAFGALILAI
						VQIIRVILEYLDQRLKAAENKFS
				1		Y\CLMTCLKCCFWCLEKFIKFL
						NRNAYIMIAIYGTNFCTSAR\NA
				1		CFLLMRNIIRVDVLDKITDFLFL
		1		1		*GKLLIVGIVGILAFFFFTHRIRI
				1		VQDTSPPLNYYWVPILTVIVGS
		1		1		YLIAHGFFQRY
15626	45994	Α	15716	I	449	FRGMGDERPHYYGKHGTPQKY
						DPTFKGPIYNRGCTDIICCVFLL
	1					LAIVGYVAVGIIAWTHGDPRKV
						IYPTDSRGEFCGQKGTKNENKP
	1					YLFYFNIVKCASPLVLAGI\QCP
}						TPQICVEKCPDRYLTYLNARSS
	1					RDFEYYKQFCVPGV/PRTIKKA
		ļ				NGVLEAGNSPCAYLKITPSLGT
		ľ				DY/PWFLLEFQCPTPQICVEKCP
1	1					DRYLTYLNARSSRDFEYYKQFC
						VPGVQEQ
15627	45995	A	15717	627	1665	EQAGQALSTAPVGDGHELLCE
l						VGSSSFRDVLGT\$MGLCGGY/G
	1					GAS/GMGGITAVTVNQNVLSPL
				į		NLEVDPNMQGMHTQEKEQVK
						TLNKSASFVDKVLFLEQQNKM
l		1		1		LETNWSLLQQQKPAQSNMDN
	ŀ	}				MFESYINNLLGQLEILGQEKLK
		1		İ		LEAELGNMQGLCMFFVKEDFK
		į.				KKYEDEINKPTEMENEFVLIKK
		ĺ				DVDDAYMNKVELEFRLEGLTD
		1				EINFLRQLYEEEIPELQSQILDTS
		1				VVLSMDNSRSLDMDSIIAEIKY
						EELQTLAGKHWDDRRLTKTKIS
		1				EMNRNISRLQAEIEGLPQRPEQ
l	1		1			RASLESVIADAEQRGELAIKDA
l	i .	1	1	1	1	KTKLSELEAAQQRAKQDMAHG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide delction, \-possible nucleotide insertion)
15628	45996	A	15718	3	3908	STQQKTARSNMDNMFESYINN
13028	43770	^	13710	ľ	3700	LRRHVETLGQEKLKLEAEL\GN
		1				MOGLVEDFKNKYEDEINKRTE
						MENEFVLIKKD\VDEAYINKVE
						LESRLEGLTDEINFLROLYEEEI
						RELOSWISDTSVVLSMDNSRSL
		1				DMNSVIAEVKAQYEEIANRSQA
1		l				EAESMYQIKDEELQSLAGKHG
		l				DNLRCTKT*DLLR*TRNISRLQA
		l				EIEGLKGORAFLEAAIADAEOC
1		l				GELAIKDANAKI/SELEAALOR
		l				AKODMA WOLR\LMN
15629	45997	A	15719	1	804	MKPRTLVVSVTVLKDGVSRVC
1		l				SFRCSDVSRISSFQWVRGLADF
1		l		ŀ		RNEATDPHGLLDDILTLRQDKG
		1				SFMICWLKGKAPPGDPAPPRITP
		1				TVRFSTMDLSAASHRIP\LSDGN
ı	İ	1			1	SIPIIGLGTYSEPKSLWATNHVP
		1				EMVRPTLERTLRVLQLDYVDL
		1		1		YIIEVPMAFKPGDEIYPRDENGK
		ļ				WLYHKSNLCATW\EAM\EACK\
		1	ŀ			DAGLVKSLGVSNFNRRQLELIL
		1		ŀ		NKPGLKHKPVSNQVECHPYFT
		1				QPKLLKFCQQHDIVITAYSPLW
						TSTNP
15630	45998	Α	15720	476	906	WKQHSHHQTWYLLTSEACKD
		l			1	AGLVKSLGVSSFNSR\QLELIQN
		l	l	İ		NPGLKPKPVSNEVECHPYYTQP
			İ	İ		KLLKFRQQHDIVMIAYSPLGTT
	ł	1		ľ		RNPTWTNMSSLPSLKDALLKR
	1	l	ļ	İ		CNKITAQVILRFNIQQGVVVIPK
		Ļ.				SFNPEKIKENLQA
15631	45999	A	15721	138	217	RHAPQVLGGLSEGT*GGHRCPA
		_	1.5000			RGHW
15632	46000	В	15722	8	578 3087	
15633	46001	A	15723	1	2232	
15634	46002	A	15724	1	2085	
15635	46003	A	15725	1	1512	
15636	46004	A	15726	1	1260	
15637	46005	A B	15728	1	2145	
15639	46006	A	15729	1	1353	
15640	46007	A	15730	1	1741	
15641	46008	A	15731	1	1995	
15642	46010	В	15732	1	3464	
15643	46010	A	15732	2	109	FTFWHDFAAAGTGCSFPCLVLP
						SWW*QNLSAFACL
15644	46012	В	15734	90	1247	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15645	46013	Α	15735	ı	1698	MELKAKAQELREECRSLRSRC
						NQLEESVSVMEDEMNEMKREG
1						KFREKRIKRNEQSLQEIWDYVK
1				1		RPNLRLTGVHESDGENGTKLEN
						TLQDIIQENFPNLARQANIQIQEI
1			1			QRTPQRYSSRRATPRHIIVRFTK
1						VEMKEKMLRAAREKGWVTHK
						GKPIRLTVDLSAETLQARREWG
						PIFNILKEKNFQPRISYPAKLSFI
						SEGEIKSFTDKQMLTDFVTTRP
				ļ		ALQELLKEALNMERKNRDTHR
						LKIKGWRKIYQANGKQKKAGV
				i		AILVSDKTDFKPTKILKDKEGH
				i		YIIVKGLIQQEELTILNTYAPSTG
						APRFIKQVRSDLQRDLDSHTLII
						GDFNTPLSTLDRSTRQKVNKDT
				ļ		QELNSALHQADLIDIYRTLHPK
1						STEYTFFSAPHHTYSKIDHIVGS
						KALLSKCKRTEIITNYLSDHSAI
		l				KLELRIKKLTQNHSTTWKLNNL
						LLNDYWVHNEMKAEIKMFFET
		l				NENKDTTYQNLWDTFKVVCTG
					İ	KFIALNAHKRKQERSKIDTLTS
				ľ		QLKELEKQEQTHSKASRRQEIT
		l				KIRAELKEIETQKTL/QKINESRS
		_				WFFERINKIDRPL
15646	46014	Α	15736	1043	1536	TSTELSTPNQQNIHFFQHHTTSI
				1		PKLTTYLEVKLSS*KKH*TWKG
Í				ļ		TTGTSCC/INHAKIVTHRLKIKG
						WRKIYQANGKQKKAGVANLV
		1				SDKTDFKPTKIKRDKEGHYIMV
		1				KGSIQQEELTTLNIYAPNTGAPR
		l				FIKQVLSDLQRDLDSHTLIMGD
						FNTPLLTLDRSTRQKVNKDTQE
						LNSALHQADLIDIYRTLHPKSTE
1						YTFFSAPHHIYSKTDHILGSKAL
1						LSKCKRTEIITNYLSDHSAIKLE
	į.					LRIKNLTKNRSTTWKLNNLLLN
				1		DYWVHNEMKAEMKMFFETNE
						NKDTTYQNLWDTFKAMCRGK
l						FIAVNAHKRKQERSKIDTLTSQ LKELEKQEQTHSKASRRQEITKI
15647	46015	A	15737	1208	1387	KPWOKNYVTHAQASVADLIK
13047	40013	^	13/3/	1208	1367	WKKGYQ*LKIK*MK*SKKRSLE
1			1			KKE*KETNKASKKYGTI
15648	46016	A	15738	308	487	KPWHKNYIMHAQASVANLINW
13048	40010	^	13/38	300	1407	KKGYQ*LKIK*VK*SEKRSLEK
1				l		
15649	46017	A	15739	220	1101	KE*KEMNKASKKYGTM
15650	46017	A	15740	1	1534	
	46019	A B		365	3904	
15651	40019	lα	15741	203	3904	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15652	46020	A	15742	79	506	EPSQRIISQDLQGWSQEHQVCL HRVGWGHCATCILPAATASHE GPEAWTARLLVPQTAVGICSA FSGLPTSTHEPTGMHFLPSEAHE KPRTQPDQKRGRDDLPCTPSTC LHTSLFLDTGQEFGTCRMAGLK EHVAGDNKERRQ
15653	46021	Α	15743	2	71	ELKF**FPILSPSRKTSPLRQGS
15654	46022	A	15744	1	627	MKABIKMFFETNENKDTTYQN LWNTFKAVRRGKFIALNAHKR KQERSKIDTLTSQLKELERQEQ THSQASRRLNQEEVESLNRPIT GSEIEAIINSLPTKKRPGPDGFTA EFYQRYKEELPKYLLKQLQOS LRIPNQCAKITSIPKHQYQTNRE PPHIETTHINCFKONKIRNPITY KGREGPLQGELRTIAQRNKRGH KQMEEHSMLM
15655	46023	A	15746	1	3828	MQLPVOTIAADGYVIKLOKRTI EDGHOSPERICIPPHLSSYSEQE TTRYPFAKAVASGFPGDTTLVF CASPQPSPQVHLPTQAVADIFSE FHEHY ARISTASVTLSKANMM GSEPTARLSNSKYPETLEVGES AQPVPMYTWKTVAODRKDG SPPLLEKQSITKSISNKPVELSSK VVEVDASKADHMKKMASMGL VVEVDASKADHMKKMASMGL VISMAGSGLVLSKSELKEISSL RNGCAIVRSEPISTDPSSWVVPR PRTNEENNGE

2008

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
15656	46024	Α	15747	ī	1639	NCLSDHSAIKLELRIKKLTONCS
						TTWKLNNLLLNDYWVHNEMK
						AEIKIFFETNENKDTTYQNLWD
		1				TFKAVCRGKFTALNAHKRKQE
		1				RSKIDTLTSQLKELEKQEQTHS
						KASRRQEITKIRAELKEIETQKT
				l		LQNINESRSLFFEKINKIDRPLA
	İ	i			ļ	RLIKKKREKDQIEAIKNDKGDIT
	i	İ			İ	TDPTEIQTTIREY\YKH\LYANKL
						EY\LEEM\DKF\LDTYTLPRLT\Q
						EEVESLNRPTTGSEIEAIINSLPT
						KK\SPGPDGFTAEFYQRYKEEL
	1		l			VPFLLKLFQSIEKEGILPNSFYE
						ASIILIPKPGRDTTKKENFRPISL
	1					MNIDAKILNKILANRIQQHIKKL
			l			IHIIDQVGFIPGMQGWFNIRKSI
						NVIQHINRAKDKNHMIISIDAG
	J		1			KAFDKIQQSFMLKALNKLDIDG
	ł		1			TAIRQEKEIKGIQLGKEEVKLSL
		1				FADDMIVYLENPIVSAQNLLKLI
	l					SNFSKVSGYKINVQKSQAFLYT
				1		NNRQTESQIMSELPFTIASKRIK
				1		YLGIQLTRDVKDLFKENYKPLL
						KEIKEDINKWKNIPCSWVGRINI
15657	46025	В	15748	1	1563	
15658	46026	Α_	15749	1	4332	

2009

15659 46027 A 15750 1 2319 MRIDGAQELTLALPFASX DVKPGPRRPLRYRCFQ LGSASLGAGFWAGRRD, DDDPSGALFSVTMDRDL DRAENIELIAQORPPRRR* GKTLQEKLYDIYVECG PQELRSNVNLLEKLVRR LLVNLYPGNQGYSVMLC SFAETIKLPYEERALLDY LPPALGDVLDKASVNIFF IVEVRDYRQSSNMQPPG ILLRITMGTLAPFVKTM* KWSQEDKFPLESQLILAT CLDPSVAVACTANRLLY MNTDPMEQCLQRYSWP, QEQSDCPPPELRVSTSG RKVGQPEELNITKAGSC* KGRPCDLAVPSEVDVEK YQSVTAADPQLPVWPAG PFRHAWEAGCOQWDTK SFNDPLLCGKIRPRKARA KSPWOFFPDDISAWLRP	KPSNR
ASFWQFFPDIBAWLEF AGRAVSQAQESVQSKVB MSHSSSG PASVSQLSSWJ PDPVWVQSSVSGKGEKE QLPSSSGKISSGNSFPPQC LKRPFPAAPAVAAAAP PAAPALAAAAVAAAAA SHSQKPSVPLIKASRRPP TRFVKIAPAIQVRTGSTG QRGGPSGPGISGSGLQ LPDARFGSSAGCPAPLCI	EQAL YSPRA KEPED SSLPC QREDG SSLPC QREDG REDG REDG REDG REDG REDG REDG REDG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
I5660	46028	Α	15751	409	1874	YDSYIVVIDDADITRGDDAVPR
						DHHSSTSMVVFRHIRLDVIVVD
						GDDQHRRMVGRRCRSAWFRR
						GSLDVDQVIGDTAAGRLNVMG
						DITRDARRDKVKNERYGVAPS
		l				VAFGLGTANRLYLNYLHVTQH
						NTPDGGIPTIGLPGYSAPSAGTA
						ALNHSGKVDTHNFYGTDSDYD
ŀ						DSTTDTATMRFEHDINDNTTIR
						NTTRWSR/S/WTWSRTANTKDV
1						SNKILTNQTNLTSTFYTGSIGHD
		l				VSTGVEFTRETQTNYGVNPVTL
ŀ		1				PAVNIYHPDSSIHPGGLTRNGA
		1				NANGQTDTFAIYAFDTLQITRD
		l	ĺ			FELNGGIRLDNYHTEYDSATAC
ĺ						GGSGRGAITCPTGVAKGSPVTT
1						VDTAKSGNLMNWKAGALYHL
ĺ				l		TENGNVYINYAVSQQPPGGNN
			ł			FALAQSGSGNSANRTDFNPQK
						ANTSEIGTKWQVRGSSRVNVM
				1		HEVNTPHQCVRFRPARSAPGCC
		l		1		QSPCHRSGKHMQYGLSKCQPG
						QYNSVKSGERVNGVTFQ
15661	46029	Α	15752	421	441	P*ARMPSPLGPPCLPVMDPETT
	1		l	ļ.		LEEPETARLRFRGFCYQEVAGP
	1	l				REALARLRELCCQWLQPEAHS
	ļ					KEQMLEMLVLEQFLGTLPTEIH
	ì					AWVRGQRPGSPEEAAALIEGLQ
						HDPGQLLG
15662	46030	Α	15753	1	346	PAFGGSAPSERLELHVDGPPPRP
						QAPGD/CGVGRSWRAEM/HDR
						ALRGTNPRRHLRSCCARARRRP
						*RRSAPPGPRAN\PS*SSWGPSTP
						GNYRCRYRSWVPHTFKSELND
						PVELLVA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15663	46031	A	15754	1	2091	MKPGLRRRRGRGLGCGLGPQL
						RATWSGAVLAGRDAVLRCEGP
						IPDVTFELLREGETKAVKTVRT
1						PGAGAKLELIFVGPQHAGNYRC
i						RYRSLVAHTFESELSDPVELLV
						AVAATLNIHLRLVHMLHIPAAG
				ļ		EGSFREGTHFLKPSPVKMPLVF
				ŀ		RVGRRRVLNQGPQGVPIPIPRSC
				l		EHLNWQKGLCGVTKCAAPLLR
			İ			GPPCPAVLRKEIPFVSLKPGLMS
				l		SKASEAEPLGPRLGSHQSTAAA
				l		LGPRYNGSADAIRPHTAWAVIQ
				1		TTISRFTWSLTQVNVPSRTVSLA
						VRRDRYMTSRTTTNPEEGCRG
1						TFWARRNGCSPSACAVAEAAL
ļ						EACLADVRYPVVALSCGRLQK
						RRDFRETNAAMSNKEKSLASA
						RFFVEVQVTLGALCTVQATGQ
						SMLVFSTTQMSSYSEKFLTSKH
						ASLSPRLQLADRWRDQGMKW
						QSPQWGELRPAASHTAEQEPGL
				ľ		AASKAFRDLQNLSNSAFLSLTS
						LSAGHQCISAIVVMMSKKEITP
1			1			QGQLAAASFGELQPVALKCPW LKSQHVESWL\RPEAHSKEQML
						EMLVLEQFLGTLPPEIQAWVRG
		١				QRPGSPEEAAALVEGLQHDPG
						QLLGWITAHVLKQEVLPAAQK
						TEEPLGSPHPSGTVESPGEGPQD
						TRIEGSVOLSCSVKEEPNVDGO
			ľ			EVAPSSPPLAAQSPEGNHGHQE
						PASTSFHPPRIQVSSPKWEV
15664	46032	A	15755	1	208	TANDEST TAT TAKE YOU TE TO THE
15665	46033	A	15756	3	373	EVAQMPRVSTPATLPSLPQPAL
1						MAPWTVPYDQLMEEEKARAR
1		l				YAGTIQKWTAAALQPLSRTSLK
						DSGEGTSKWGRSVDTTQPLSPA
						IPVIFQ*AHEQSGHSGMDGGYA
						WALQHGLALTNTDLA
15666	46034	Α	15757	3	266	
15667	46035	Α	15758	3	587	YPASAGLMLQNFGVIGLRYHF
						AIHSPAAGGLLDGLHAVAAIQG
						ITKIETTPNHQRAPAHWTLLTQ
						QAHLQPSPLHFNLPLTLCLMHC
				l		PTAIPHCFADARTWVNLPTSSLI
				l	1	GHKKENLKEFISGSLIVHEILEE
1				l	1	VLQAEGDFQPFTRVTVHWGKG
1				1		NDQTFRGLLDTGSELTLIPGDP
				l		KHHYGPPVKVG\AYGAQLL
15668	46036	Α	15759	1	1566	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
l				sequence		
		L				
15669	46037	A	15760	I	981	MVAIAKVLIAYKEDSDSHNQPN
i				l		TRMTVSKGPVSAVEGGRRGQD
						SDDCVEENDAVRTWPPSRILRT
1						ESHPSPHTCSWRRHLLSSCPYT
1						GMSFPIPSSGNISSTHAVISFPPL
				l		SEEMNPMLPKATVMTSLEAAA
						RQNLKSRQAPRCEVESVTHEEV
						YYTRKELFEFSNLYRKKSGEQI
		ŀ				REWILRKTDGSWRITVDYRKL
1	l					NOMVTPIAAAVPDV/VVSLLEO
	ŀ			1		INTSPGTW\LCHNLIWRDLDCFL
				l		LPONITLVHYTDDIMQIGSSEQE
İ		l				VANTLDLLPALMASW/ENSP*SI
	i		ŀ	Ì		DRGRED*SLVHKWFCTICRHHP
	l		i			KVDSCSTTAPF*DIPEGQQL
15670	46038	A	15761	349	718	AGPEGTTTAECP/I/COORPILS
13070	40036	l^	13701	1777	/10	LRYGTISWG/DOSATWWQVDY
			į	}		IRTLLSWKWOSASAKTTIHGLT
						KCLIHHDIPHSIASD*GTCFMAK
						EVWQWYCFSHSQDSRVQESRG
				ł		
14(7)	46000		147/0	140	400	GIGSCTTHHHPCSFPN
15671	46039	Α	15762	143	469	ARIQ/GSRNQGVEVEVAPLTVT
		l	l	1		PSDPLANVLLPVPATLPSAGLEI
						LVPEEGRLPPGDTTMMPLNWN
1		l			i	LRLPHGHFGLLLPLNQQAKKG
		_				VAVLGGVIALDCQDEISLLLYK
15672	46040	Α	15763	I	2712	
15673	46041	A	15764	3	578	LKSRQAPRCEVESVTHEEVYYT
	1					RKELFEFSNLYRKKSGEQIREWI
		1				LRKTDGSWRITVDYRKLNQMV
		l				TPIAAAVPDV/VVSLLEQINTSP
1						GTWLCHNLIWRDLDCFLLPQN
				1		ITLVHYTDDIMQIGSSEQEVAN
		l		l		TLDLLPALMASW/ENSP*SIDRG
		l		1		RED*SLVHKWFCTICRHHPKVD
				1		SCSTTAPF*DIPEGQQL
15674	46042	Α	15765	1	461	MTVDYRKFNOVVTPMAA/AVP
	1			I		DAVSLLEOINTFPGTWYAAIDL
		1		1		ANAFFSIPVHEAHOKQFAF\LPQ
	1	1		1	1	GYINFPALCHNLIRRELDFFLLL
1				1	1	QDITLVHYIDDILLIGSSEQEVV
1		1	1	I	1	NTLDLLIHKRSKEAEHTAASRIR
				1		VSCLPEOKSHEOTLPWEQVP
15675	46043	A	15766		1371	VSCLIEGRAMEQTERWEQVE
		_		54	864	
15676	46044	В	15767	124	004	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /-possible nucleotide deletion, \=possible nucleotide insertion)
15677	46045	Α	15768	2	1148	PRGRNRRRKTFQERRMTLNESP
				i		EKIGKWIECYGHPPASKLVEIYI
						HTVFVEDKLSICIRSFNKKADGS
						WRMTVDYCKLNQVVTAIAAAI
					Į.	P\EWFHCLSKLIHLLIPGMQPLT
ļ.						WQ/HALFSIPVHKGYINSLALC/
						Q*CYLERT*LLFTSTRYHTGPLH
						**HYADWIQCARSRKQTGLTG
						KGQRFVLTERDTY\YGFTYPAH
				ł		NASAKTTIHGCTECLTHPHVIPH
						SIASEQDTHFTAKEVWQWAHG
						NGIHWSYHVPHHPEIARLIEWW
						NGLLKSQVQCHLGDNTLQGW
						GKVLQKAVYALNQYPIYGTVS
				İ		PIARIHGSRNQGVEVEVAPLTIT
						PSDPLAKCLFPVPKTLCAACLE
1						VLLPGGGMLPPGDTTTIPLNRK
						LRFPLEHFGLL
15678	46046	Α	15769	917	2352	DTISHQLEWRSLKSQETTGIAEI
		l				SATIKDLKGAGVVIPTTSLFNSS
						VWPVQKTDKSWRMTVDYCKL
1						NQVLTPNAAAVPDVVPLLEQIN
						TSPGTWYAAIDLANAIFSIPVHK
						AQQKQFAFSWQGQQYTFTVLP
			İ			QGYISSLALCHNLIRRDLDGFLL
						LQNITLLVHYVDDIMLIGSSEQE
		1			1	VANALDLLVRHLCVRGWEISPT
1	1	1				KIQGPSTLVKFLGVQWCGACR
						DILSKVKDKLLHLAPPTTKKEA
1					1	QRLVGLFGFWRQHIPHLGGLLQ
1	ļ					PLYLVIQKAAIFEWNPEPEKVL
		l		ļ		QKVQAAVQAALPLGPYDPTDPI
		l				MLEVLVADRDAVWSLWQALI
		l				GESQQRPVGFWSMALPSSADN
				i		YSPFERQLFACSWALVETECLT
				l		MGHQVTMQPELPIMNWMLSDP
		1		i		SSHKVVQAQQH\QEVAQMPRV
		l				STPATLPSLPQPALMAPWTVPY
						DQLMEEEKARARYAGTIQKWT
10070	16045	ļ	15770		861	AAALQPLSRTSLKDSGEGTSKW
15679 15680	46047	A	15770	5	90	
15681	46049	A	15772	1	324	MVQEASEVIGQLQSSAAK*AM
13001	70047	ľ	13/12	ľ	727	HWAGFSYLTKKSLNAN
15682	46050	С	15773	1	726	
15683	46051	Ā	15774	364	543	ARGAARSMKHTSGDETKRQKL
1		1				D*VRLTGIRASREEVTKEAGPR
		l		l		KRIMCSGMDELNGGLQL
15684	46052	В	15775	1	1056	
15685	46053	A	15776	1	282	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, *-possible nucleotide insertion)
15686	46054	A	15777	167	954	IKNTSKLESPRGSSPHCSRNPV LLRGIGRYSRSAMYSKAMYK, RKYSAAKSKYEKKKKEK VLAT VTKPVGGDKNGGTEVVKLRK PSQHVKKLRASITFGTILIILTG PLVLNRIVPLRRTHQKFVILLTG PLVLNRIVPLRRTHQKFVILATFI KIDISNVKIPKHLTDAYFKKK LEKPRHQGGEIPTEKEKYEITE QRKIDQKAVDSQLEPKIKAIPQL GOYLBSVEALTNGIVPIKIQL
15687	46055	A	15778	1054	1403	MVEGGILRVNTSVECKTGPFFF SPHILWREPFCSVQVGKKKKEK VLATVTKPVGGDKNGGTRVVK LRKMPRYYPTEDVPRKLLSHG KKPFSQHVRKLRPSITPG\TILIIL TGNRHG
15688	46056	Α	15779	1	388	
15689	46057	A	15780	1	456	MKLPEEGSLSDICCSAIFAVLQP LLVIPRETGSGVDLQQTPTDLQ LRQTESQIMSELPFTVASKRIKY LGIQLTKDVKDLFKEIYKPLLSE IKEDTNKWKSIPCSWLGRINIVK MAILPKVTYIFNAIHIKLPMTEL EKTKHIWNKKACI
15690	46058	Α	15781	1	504	
15691	46059	A	15782	1	1836	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
15692	46060	Α	15783	285	1949	RGEMNSRGRNNLSIINKTDFPFI
						ALNAHKRKHERSKIDTLTSQLK
						ELEKQEQTNSKASRSHEITKIRA
	ì					ELKEIETHNKPFKKISESR\SWFF
	1					IFYF\ERINKIDRPLARLIKKKRE
						KNQIDTIKNDKGDITTDPTEIQT
	l	i		ŀ		TIREYYKHLYANKLENLEEMD
						KFLDTYTLPRLNQEEVESLNRPI
	1					TGSEIEAIINSLPTKRSPGPDGFT
				ŀ		AEFYQRLKQCRKAHLDPVSLT
l						HRSGPGAREDRLSDSGDASRGC
				ļ		WNLSQHAAVTSPESEERRIRTL
l						SCRSVGVCRRSTPDPVCLGISSG
1	1					GCRTVNIGEQQMLLPDRSSGSF
				1		VSEGYRPCEVSAPTGGCPPVRC
ŀ	1					LMINTVSLIGVKFTEVFDSINISE
						HLELACLQDHNSLLAREQNWT
	1					EDEFDELTEVGFRRWVITNSSE
						LKEEVQTHCKEVKNLETRLDE
		l			ĺ	WLTRITNAEKSLKDLMELKTM
						AGELRDACTRFSSRFNQLEERV
l		1		l		TEIEDQMNEMKQEEKFREKRIK
		1		[ĺ	RNEQSLQEIWDYVKRPNLHWL
1				ŀ		GVPESDRENGTKLENTLQDIIQE
						NFPNLARQTNIQIQEMQRMP
15693	46061	В	15784	596	1414	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15694	46062	A	15785	610	2986	QTERNSININRNGHQHQNPICRS
						PSSKTKGSIKPQRSGRNQSRKA
i	ł				ļ	ENSKDORTSSPTKDCSSSPATE
į .				1		ORWMENDFDELTEV/VLOKPG
	ŀ					RDTT\KQKEKFRPISLMNIDVKI
i						LNKILANRIQQHIKKLIHHDQV
						GFISGMOGWFNICKSINVIHHIN
	ŀ					RTNDKNHMIISIDAEKAFDKIQ
	i					QPFMLKTLKKLGIDGTYLKIIRA
	ĺ					IYDKSTASIILNGQKLEAFPLKD
	ŀ				İ	RTRQGCPLSPLLFNIALEVLARA
	ĺ					IRQEKDIKCIQLGKEKVRLSLFA
	ŀ					EDMIVYLENPIVSAPNLFKLISN
	l					FSKVSGYKINVQKSQVFLYINN
	l				1	RORESQIMNEFPFTIARRRIKYL
1	ľ		1			GIOLTROVKOLFKENYKPLLKE
						IKEDTNKWKNMPCSWIGRINIM
			ĺ			KMAILAKVIYRFNAIPIKLPMTF
						FTELEK\TTLKCI*NQKRAHIAK
	1					PILSOKNKAGGITLPDFKLHYK
			ĺ			AAVTKTAWYWYONRDIDORN
	ŀ		İ			RTEPTEIMLHIYNHLIFDKPDKN
						KKWGNDSLFNKWCWENWLAI
						CRKLKRDPFLTPYTKINSRWIE
						DLNIRPETIKTLEENLGNTIQDIA
						MGKDFMSKTPKAMATKAKIDK
				i		WDLIKLKNFCTAKETTIRVNRQ
				İ		PTEWEKIFAIYSSDKGIISRIYNE
	ĺ					LKQIYKKK\NNPIKKWAKDINR
1	ĺ					HFSKEDIQAASKHVKKHSTSRIF
	1					MQIKTTLGYHLTPVRMATIKKS
	i					KNNRCWCGCGEKGTLIHCW/M
15695	46063	A	15786	1	906	MVNQKRYHIPGGTAEISGTIKK
					i	LKNAGVVIPTTSPLNSPIWPVQ
						KTDGSWRMTMDYHKLNQVVT
				1		PIAVAVPYVVALLEQINTFSCT
i						WYAAIDLANTFFSFSVHKA/HK
1				ĺ		K*FAFSRQGQQYTFTVHP\YDN
						YSPFERQLLACYWASVETEHLT
1			1	l		MGYQVTIQLELPIMNWVLSDPS
1						SHKLGHAKQHSIIKWKWYIRD
						RARAGPEGTARIHRSRNQGVEV
						EVAPLTITPSDPLARFLLPVPTIL
	1					HSAGLEVLVPEGGTLPLGDTTT
						IPFNWKLRLPPGHFRLLLPLNQ
1						QAKKGVTVFAGVIEAIKMK
						·

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15696	46064	Α	15787	1	909	MVLEVSVSDRDAVWRLWRAPI
						GESQQRSLGFWSKVLPYSADN
				i		YFPFERQLLACYWALLETDRLT
						VGHQVTLQPELPIMNWVLSDPS
						SHKVGHVHQHSIIKWKWYIRD
						QTRAGPEGTTTPVITQ/WDAHE
						QSGLSGRDG\KGQRFVLTGVDT
i						YSGYWFAYPAHNASAKTSIYG
						FTECLIHCHGIPHSIASDQGTLFT
				l		AKEVWQWAHAHGIHWSYHIP
						HHPIAAGLIEWWNGLLKSQLQ
		Į .				CQLGDNTLQGWGKDLQKAMY
				l		SLNQRLIYSTVSPISRIHGSRNQ
						RVEVEVAPLTITLSDPLAKFFFL
15697	46065	Α	15788	I	754	MQNNVDSFQEPPPTPLFSSRPIA
						RLKSKQAPRGEAESVTHEELESI
	l					RLRAKTSDKISSLTFRRGWGKV
					l	LQKAMYALNQHPIYGTVFPIPRI
						DGSRNQGVEVKVAPLTITLSDP
			1			LAKFLLPVLSALHSAGLEVLVP
			1			EGGILPPGAGDMRMIQLNWKL
						SLPPGYFGLLLPLSQQTKKGVT
						DLGGVIDLDYQDEISLLFHI\KV
			1	l		NGKLQQPNPDRTTSGPDPSGM
						KVWVTPPGK/EPQPAEGLAEGK
	10000		1.5000		1506	GNTEWVVEE
15698	46066	Α	15789	686	1506	VETVHLMLQIGEFKKTDVS/WK
						TVDYHKLNQVVTPTAAAVPDV VSLLEQINTYPGTWYAVIDLEN
			1			
					1	AFFSVPVHNAHQKGFAFSWQG QQYTFTVLPQGYINSPTLCHNLI
		l				
		i				RRDPGHFSYPQDITLVHYIDDI MLIGSSEKEVANTLDLLPIYOV
			l	i		TLKAANYEWGPEQEKAVQQV OAAVOAALPLGOHNPADPTVL
				i		
				l		EVSVADRDAIWNLWQAPIARIH
		1				GSRNQGVEVEVAPLTITPSDPL
			l			AKFLPPVPATFCSAGLEVLVPE
15500	14047	<u> </u>	15700	400	507	GGMLPPGEIETIPLK
15699	46067	A	15790	422	537	VNQFGVIFCAAIIN*YECLKQSF
15000	15050	<u>_</u>	15701		510	KTQTDYFPSPLNTSQ
15700	46068	В	15791	1	519	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		<u> </u>		ļ		
15701	46069	Α	15792	53	753	SLGTGTLRVLLGTAEAQMLQFS
i						HLYADYGGKPALEDINLTLESG
		l				ELLVVLG/PSGCGKTTLLNLIAG
1		l			1	FVPYQHGSIQLAGKRIEGPGAE
		l				RGVVFQNEGLLPWRNVQDTVA
	1	1				FGLQLAGIKKMQRLKFAHQML
						KKVGLEGADKRYIWQLSGGQR
						QRVGIARALAANPQLLLLDEPF
		l	ł			GALDAFTRDQMQTLLLNLWQE
		i			1	TGKOVLLITHDIEEAVFMATEL
						VLLSSAP\GRVLERLPHI
15702	46070	A	15793	2348	2760	LWLVCINCFIPAFLLMLRWLPR
						LKPLSQLSVPTYRLPSVPFLKPT
				į.		GMESPEAISRVSGKGIL*RDSDA
					1	YSVSLSYHYPIIRHPSSIIVDHRS
1						SIDP*PIVDHSSSIAKRSGTFPLG
		l	1			YWEVFVQLTFGQQRQASGILQ
15703	46071	A	15794	2	415	HPGLETTDLLSPLPAALEPQGQ
13703	40071	l^	13794	-	1713	GRPSIQGAPRGSFTRIEAGTPHV
					1	PAPCPAAORNOGWGLVGAWL
			ĺ			GFMAP*EOSVASSWK*PLMKS
						QYSREVISHLPLVPMQQGRPVM
1						TOSMWGWLVWFPALFLVOIGH
1					1	
15004	44070		15705	400	005	LVMSSGHS
15704	46072	A	15795	428	885	GISMKLSRISAI/NWNKISDDKD
		i				LEVWNR/LTSNFWLPEKVPLSN
		l				DIGAWQTLTVVEQQLTMRVFT
					i	GLTLVDTLQNVIGAPSLMPDAL
1		ĺ				TPHEEAGLSNISFMEAGHARSY
I		ŀ			ı	SSIFSTLCQTKDVDAAYAWSEE
		l				NAPLQRKAQIIQQHYRGDDPLK
		L_				K
15705	46073	Α	15796	1	1125	
15706	46074	Α	15797	57	303	SHKHYADALLYPAPDVCTAC*
		l	1			LLRGNRRDQSKS\TAKRHLTAA
		1				HFRFVRMVRHGNH/SRLRPDSF
		<u></u>				R*PVSPDR*PPLRAAHFR
15707	46075	A	15798	3	430	
15708	46076	В	15799	I	1338	
15709	46077	Α	15800	380	1043	WTRWRASAADNLAHGRWGEQ
		1	l			KLVVTTGASGAPDRQAARCGC
		1	1			IWRKISWYRSKGSKHPLTPKEV
1		1				EKTGWGLKIRGMGKQQIA\ERV
1		l				KEAARILELDGLLKRRPRELSG
		I				GQRQRVAMARA1VRDPAVFLF
1		I				DEPLSNLDAKLRVOMRLELOO
		I				LHRRDQVEAMTLAQRVMVMN
1	1	1				GGVAEOIGTPVEVDGDDTLEIL
1	1	1	1	1		GADNLAHDAGRAEAGGDWRI
	1	1		1		RSARRSSTCAH
			L		1	TO LOCALI

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15710	46078	A	15801	1	900	MAAIAFLITWFLSHDKKRIRILK
				ŀ		RLSGGGNMADEFPGGAVVFTV
						LNLTHWLEDWGLPVHWRSSC
		l		i		MPSGLEYLIVLIFKGVIVAVHLL
i				ł		IVDALNLIRRIHAV/QGSPCVET
	l					CQHALDQLIMHSQPTHAVAVF
						DDENRSSGWRHQGLPDYKAGR
	1					PPMPEELHDEMPALRAAFE\QR
		ĺ				GVPCWSTSGNEADDLAATLAV
						KVTQAGHQATIVSTDKGYCQL
1	1					LSPTLRIRDYFQKRWLDAPFID
l	1					KEFGVQPQQLPDYWGLAGISSS
	1					KVPGVAGIGPKSATQLLVEFQS
						LDGIYENL\D\AVAEKWRHKL
15711	46079	A	15802	1	1350	
15712	46080	Α	15803	1618	2082	
15713	46081	A	15804	1	803	MINLIGSDVNYDWLKLPLVHL
		1	I	ł		HWYDKEVRPGRKVGHLNLTDS
		1				DTSRLTATLEALIPLLPPEYASG
		1				VIWAQSLIRCVKHRIRHCAPIAG
Ì		1				CGTGCRPDKTRQASHQAQMSN
						AYDYSEIQGIGKQKAANRVAEP
		1	l			RTYHSFASITDRRADLLKPGNH
						AQDGRLTAARRADKHDELFIG
					İ	NFQKADNRGQVSDSVEGNSNC
						APVKPAPIPTERIPLALMASMPA
						CASSMPIHACGGTPSPAQAARK
						ISGSGFEFLSWEASAQKVKNSR
		1		Į.		SSSAAKIVGAFRETEASPTEICC
				[ARRSASKERMPG*ARADPHRA
	1			[HTAGVNGVNARLRIFNADTCL
			l	1		RRHAQPCTGGEKDFRVRFRVF
		1				KLGSIRAEGEKLAQLQRR

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SEQ ID	SEO ID NO	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1.00	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequence		
15714	46082	A	15805	1	1655	MRGHNWIDMDQARQQSTPHT
13/14	40082	l ^A	13803	1'	1033	DRKICHKNDRPRTTPLRKIDAH
		1				PHTLIILPAESTLSTTKYKLRAL
					ļ	LVENTNKDKSAVSVVFRGSVL
		1				
		1				HQQAANFPNLMLCFQLKNGVN
				l		ACNOFIMLEVODONAAGSIERAA
1		1		l .		SPCINEKQHAFSESVPAIAAVRG
				ı	i	LGSMIAVEFNDPQTGEPSAAIA
						QKIQQRALAQGLLLLTCGAYG
1		1				NACKRRIDYVCRIKHSRRIRQV VLLNFAKSGAFSTTRGTDDKLT
1						
				1	i	PVIQPRASGEYNEEFSLLPVVN
1		1		1	1	YLKDKLSNPVRLVKDYLDGVD
		1			l	VAEGELVVLENVRFNKGEKKD
						DETLSKKYAALCDVFVMDAFG
1						TAHLRAGFLLTGI\GKLRCSLRC
I		1				AGPLLAAELDALGKALKEPARF
		1		1		MVAIVGGSKVSTKLTVLDSLSK
		1		i		IADQLIVGGGIANTFIAAQGHD
		1				VGKSLYEADLVDEAKRLLTTC
				I		NIPVPSDVRVATEFSETAPANL
1				ì		KSVNDVKADEQILDIVRDLDAA
1			1	ł		NVVINAVFSYVTNVWGWAFE
			1			WYMVVMLFGWFWLVFGPYAK
16716	46002	١.	15906	ļ	1359	KRLGNEPPEFSTASWIFMMFAL
15715 15716	46083	A	15806	670	1008	
15717	46085	A	15808	455	1137	GTKPRDIDHILGTFITPGMPKGG
13717	10000	ľ	13000	1.55	1	KLDVYAAPELPLKLLGRPTEGE
1						YNEEF/SLLPVVNYLKDKLSNP
						V/RLDGVDVAEGEL/VVLENVR
				l .	į.	FNKGEKKDDE/HRAQASTHGIG
			i			K/FADVACAGPLLA/AVLDSLSK
		1			1	/IADQLIVGGGIANTFI/AAQGHD
						VGKSLYEADLV/DEAKRLLTTC
						NIPVPSD/VRVATEFSETAPATL
		1				KSVRHSDKISYISTGGGAFLEL
		1				WKVKYLPAVAMLEERG
16710	46086	A	15809	465	888	RGLVALSSERRCRLRRRNHPES
15718	40080	l ^A	13809	703	000	GTRSPDPAGAASLSRRFFYAPP
		1	1	I		
1				1		GTPETGCGYSSRSRSQRKR/LST WRCNS*ETPKSPPKACCRPARIP
				1	1	APRSSSVKKASACGPAAVMKK
1		1		1	1	RCRKASITPISKITCAIHRMRRW
				1	1	
1.0016	46007	١.	16010	417	707	TCTKRAAF
15719	46087	Α	15810	417	707	IQIHERSWRYLHWWSSVRRRA
1		1		1		A*YGAPL*RDGIQRGTRRISQV
1				1		YQ*SSVRQLRRDASASGSAPRV
					[CREYAAACYLRHEHWSCDAVS
	46000	1	1.0011	 	1250	PTPRMLKSFV
15720	46088	В	15811	1	4350	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15721	46089	В	15812	1	2862	
15722	46090	A	15813	1	1471	MLHSRGFLAEVFGILARHNISV
						DLITTSEVSVALTLDTTGSTSTG
		l				DTLLTQSLLMELSALCRVEVEE
						GLALVALIGNDLSKACGVGKE
		l				VFGVLEPFNIRMICYGASSHNL
		1		1		CFLVPGEDAEQDGTGTSGIGAQ
		l				KKKMYANNGAIDRKLLFEATF
		ı				VTIEKCCDTNQKGDDTHALGQ
		1				PIRGHDKSLAGSFCYACRSEEG
		1				LSQYRAYDSRGQLIAVKDTQG
	ŀ					HETRYEYNIAGDLTAVIAPDGS
						RNGTQYDAWGKAVRTTQGG\L
						TRSMEYD\AAGRVIRLTSENGS
	l .					HTTFRYDVLDRLIQETGFDGRT
	i	1				QRYHHDLTGKLIRSEDEGLVTH
	i					WHYDEADRLTHRTVKGETAER
		1				WQYDERGWLTDISHISEGHRV
	i	1				AVHYRYDEKGRLTGEROTVHH
	i	l			l	POTEALLWOHETRHAYNAOGL
		l				ANRCIPDSLPAVEWLTYGSGYL
		l	İ		1	AGMKLGDTPLVEYTRDRLHRE
		l	İ			TLRSFGRYELTTAYTPAGQLQS
		1			}	QHLNSLLTYRHANFAL
15723	46091	Α	15814	1	1367	MSKLLERFRYFKQKGETFADG
					1	HGQVMHSNRDWEDSYRQRWQ
		l				FDKIVRSTHGVNCTGSCSWKIY
			İ			VKNGLVTWEIQQTDYPRTRPD
		l			İ	LPNHEPRGCPRGASYSWYLYSA
		l				NRLKYPLIRKRLIEL WREALKQ
	1	l			l	HSDPAADNDLIARFQSFTNQPFI
		l				ANRLTGNDRARFNLTFIVNNQH
		l				ATAITTTGTFLPEWSTTFIITLIL
		l	1			QSKVQAIDLTKEAIVTVGCNTIS
						TTVHEVVSASIGIATKFWQHIGF
		l				CFYVIYDTVVTTIVESRTTEYSP
		ł				DVVTGLITRITTPDGRASAFYN
		l				NHHSQLTSATGPDGLEIRREYD
	1	1	ĺ	l		EWGRLIQETAPDGDITRYRYDN
	1	1	ŀ			PHSDLPCATEDATGSRKTMTW
	1		İ	l		SRYGQLLSFTDCSGYVTRYDH
	1	1	l			DRFGQVTAVHREEGLSQYRAY
	1	1		1		DSRGQLIAVKDTQGHETRYEY
		1				NAA/GDIRQCESDLQSPSCSQYR
	1	1		l	1	AIDGESLSETARYSVHO

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \"possible nucleotide insertion)
15724	46092	Α	15815	357	1733	TADSPETLRCQTSGENARGAGT
						VERAGKSLWRSLR*SHRP*STR
	l	ŀ				TGDIHCQIQTNFSLFN*QHPGQH
						R\YGSGTAIINQSLITHDPEYPEN
						LPAAPLVRYGWTPRGELAVVY
						DRSGKQVRSFTYDDKYRGRMV
1						AHRHTGRPEIRYRYDSDGRVTE
					8	QLNPAGLSYTYQYEKDRITITD
						SLDRREVLHTQGEAGLKRVVK
						KEHADGSVTQSQFDAVGRLRA
		1				QTDAAGRTTEYSPDVVTGLITR
						ITTPDGRASAFYYNHHNQLTSA
		ĺ				TGPDGLELRREYDELGRLIQET
						APDGDITRYRYDNPHSDLPCAT
						EDATGSRKTMTWSRYGQLLSF
	ľ					TDCSGYVTRYDHDRFGOMTAV
1				1		HREERLSQYRAFDRRGTVIAVK
1		l				ETOGLETGFEYHFPVPDGVIAP
1				Ì		KG/SRNGNSQSVGKRRSTEDIR
1		l				G*TRWRGYTADRRAAKTHARV
		1				E*HLRARPRVCALFFPKQCERH
15725	46093	A	15816	1595	3228	DNNAKLFRAGOMKVLKPYVDS
		1		1		GKIKVVGDQWVDGWLPENAL
		1		l		KIMENALTANNNKIDAVVASN
		1		1	1	DATAGGAIQALSAQGLSGKVAI
		ŀ		l		SGODADLAGIKRIAAGTOTMT
		ŀ				VYKPITLLANTAAEIAVELGNG
		ŀ	i e			QEPKADTTLNNGLKDVPSRLLT
		ŀ		l		PIDVNKNNIKDTNLPAAPLVRY
		ŀ		l		GWTPRGELAVVYDRSGKQVRT
İ		1				FTYDDKYRGRMVAHRHTGRPE
1		l				IRYRYDSDGRVTEOLNPAGLSY
				l		TYOYEKDRITITDSLDRREVLH
i						TQGEAGLKRVVKKEHADGSVT
1				l		QSQFDAVGRLRAQTDAAGRTT
						EYSPDVVTGLITRITTPDGRASA
1		1				FYYNHHNOLTSATGPDGLELR
		l	l			REYDELGRLIQETAPDGDITRY
1		1	l	l		RYDNPHSDLPCATEDATGSRKT
1			1	1		MTWSRYGQLLSFTDCSGYVTR
1	1	1		i		YDHDRFGOMTAVHR\EEGL\SO
1		1				YRAYDSRGQLIAVKDTQGHET
				l		RYEYNIAGDLTAVIAPDGSRNG
1		1				TQYDAWGKAVRTTQGGLTRS
		1		1		MEYDAAGRVIRLTSENGSHTTF
				1		RYDVLEQYCGQVIKAHAVSRF
15726	46094	A	15817	1	627	KIDI DEQICOQVIKANAVSKI
	46095	A	15818	701	895	
15727	40093	I ^A	12919	/01	07.7	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15728	46096	A	15819	1412	1705	LASISTSCAPPKVNCGVMAIFLA
		1				SDSRLWSGLRAQATSRSTLSLM
					l	TRSGLWLAIO*NALRSWLASHE
						TKRPPSTLGGNTVPS*ALRLGIL
						ASSRILL
15729	46097	Α	15820	344	547	
15730	46098	Α	15821	1	1518 '	
15731	46099	Α	15822	1693	2455	VLAAFTRSKQRYGAPRLTDELR
l						AQGYPFNVKTVAASMRRQGLR
						AKASRKFSPVSYRAHGLPVSEN
1		İ				LLEQDFYASSPNQKWAGDITYL
İ						RTDEGWLYLAVVIDLWSRAMA
						LLRRKSPGNVIVHTDRGGOYCS
						ADYOAOLKRHNLRGSMSAKG
	ļ.			l		CCYDNACVESFFHSLKVECIHG
						EHFISREIMRATVF\IISNVITIGG
						GGTVAARGGVP*AVKSNVNDV
				ŀ		RRNCLARSSNNGWIFPPVVVNG
l						WROOALTIVAGICC
15732	46100	В	15823	1	1407	
15733	46101	Α	15824	1	3252	
15734	46102	Α	15825	1	2409	
15735	46103	В	15826	1	2802	
15736	46104	Α	15827	1715	1824	
15737	46105	Α	15828	410	720	PGSRAVIGWSMSPRMTAHWPA
						MPCRWRCGGVRTYAPEPTG*R
	ŀ					RWLQNCSACSS**LRKTSRKNR
1		1				TT*RTPARSSSAPPLMLKQPFTR
		1				QWPLNTPWKPPRLSPS
15738	46106	Α	15829	905	1008	SCDSYRLSSRAAAVTPMRSASF
						RASGFLLVILFVCRVSVVGCRV
						LAWSGGSVFRRGVPLAWTVLR
			l	l		WVRRRCGCGVGRCRGLSAVV
	1			l		YPSGIITVDSARSWRIMGRSGSP
l				1		LA\TVGLAGCWSOPRL*TDAFS
l				l		KLOGFRVFTGDTICM
15739	46107	В	15830	ī	3148	
15740	46108	Α	15831	1896	2062	SVCYPGRN*IPSG*N*PRIWSDP
1			1	l		VQRRERAGKKWLHENPGWRK
				1		**PDYHYRSEAD

SEQ ID NO:	SEQ ID NO: of peptide sequence		in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
15741	46109	A	15832	1	1456	MSTEIARLKRQLADGMKSWLS SKRPRHTSRAPEMKYVFIEKHQ AEFASKQCAACSGWPATAGNV VSAADKDKHASAVPPHCDSVV LAAFTRSKQRLPLNVKTVAASL RRQGLRARPFGSSARSATAHTA CLVRKSVGAGFLRQWPEPKWA GDTTYLKTDEGWLYLAVVIDL WSRAVIGWSMSPRNTAQLACD LOGALGMALWRKFRENVIVIDL GGQFGDRWKALANALNIPLDQ LSARISANPKGGFUYLAREVNP MADYIKKLKPGHVYRESSR YYPSGEVTAHLIGFTNVDSQGI EGYGKSFGKWTGACGEGWKTGACGKWTG DRYGRVIEDISSTDSQAAHNLA LSIDERLQALVYRELNNAVAFN MSPSYNPNNLSGTPKEAMRNRT LDVFERGSTVKPMVWMTGLOR
						GVVRENSVLNTIPYRINGHEIKD VARITTEEDFNHASAARFVCAA AERALONHH
15742	46110	A	15833	2	929	
	46111	A	15834	1	472	
	46112	A	15835	1	513	
15745	46113	A	15836	ı	254	CDNLKTCHTSHGSVMAETAVI NHKKRKNSPRIVQSNDLTGAA YSLSRDQKRV/LRYEPVSQFRLS ETQKHQSVCHAFIRIPVSVS

2025

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.01	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
15746	46114	Α	15837	145	1929	VSGVIEIADGSRRRKAAALTES
						DYRVLVGELDDEQMAALSRLG
						NDYRPTSAYERGQRYASRLQN
ŀ						EFAGNISALADAECDNLKTCHT
						SHGSVMAETAVINHKKRKNSP
						RIVQSNDLTEAAYSLSRDQKRM
ł						LYLFVDQIRKSDGTLQEHDGIC
1						EIHVAKYAEIFGLTSAEASKDIR
						QALKSFAGKEVVFYRPEEDAG
1			ŀ			DEKGYESFPWFIK/RPSRGLYSV
	l			ŀ		HINPYLIPFFIG\LQNRFTQFRLS
				l .		ETKEITNPYAMRLYESLCQYRK
						PDGSGIVSLKIDWHERYQLPQS
1				1		YORMPDFRRRFLOGFCRFRNH
1				ļ.		HQTGFSPAGANQRGPLAATLSG
					1	PGGEGQSAVARLTGEKKNHPG
i i						AQYANRLSPRVGRFINAAGTTG
l]	İ				FPTWKAGSERNAINDDVTYAIK
ı				l		PTCWPGLDIIPSCLALHRIETEL
1				l		MGKFDEGKLPTDPHLMLRLAIE
1				l		TVAHDYDVIVIDSAPNLGIGTIN
1			1	İ		VVCAADVLIVPTPAELFDYTSA
				l		LOFFDMLRDLLKNVDLKGNSN
1						GSQSPWMEEQIRDAWGSMVLK
				İ		NVVRETDEVGKGQIRMRTVFE
				1		QAIDQRSSTGAWRNALSIWEPE
				i		CNEISIGVSLDQDGGSNSVLRK
15747	461I5	В	15838	I	1734	
15748	46116	Α	15839	1	97I	
15749	46117	Α	15840	1	1143	
15750	46118	Α	15841	1	556	
15751	46119	В	15842	8	1000	
15752	46120	Α	15843	1	1169	MLTLILLDLQFMAQLAGYQMS
				ĺ		FNSTKSKQTKNNFAVGYRTGD
				l		FQLHTNVNDGAEFGGSVYQKV
						CEELDALINLAWTSSTSCTRFG
						LAAKFQLKPIAFISTKVNHWLT
						GVSYTPPLEAGVKLNLSALCDN
				ŀ		LKTCHTSHGSVMAETAVINHK
						KRKNSPRIVQSNDLTEAAYSLS
				i .		RDQKRMLYLFVDQIRKSDGTL
				ļ		QEHDGICEIHVAKYAEIFGLTSA
						EASKDIRQALKSFAGKEVVFYR
				1		PEEDAGDEKGYESFPWFIKRAH
			1			SPSRGLYSVHINPYLIPFFIGLQN
1						RFTOFRLSETKEITNPYAMRLY
1			l			ESLCQYRKP\DGSGIVSLKIDWII
						ERYOLPO/TYORMPDFRRRFLO
		1	l	l		EEFESLNRQITGSEIVAMINSLST
						KKSPGPDGFTAEFYQ
15753	46121	Ā	15844	1	1509	

NO:	SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
15754 46122							
15754 46122				09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
15755 46123	1	1			sequence		
15755 46123	15754	46122	_	15945		1559	
15756 46124			-				
15757 46125							
WISAGAAPGGPIVKQLPSPVRT							MENT LOWLETT COTTOUR
EQLEQERPYTIGS.APSTPNIKLLL SHSNCQLPRPLRSITNISACFSLE LAIDHRGPPHSPCARSLYSNSVY	15757	46125	A	15848	11	812	
SHSNCQLPRPLRSITNISACFSLE LAIDHRGPPHSPGARSLYSNSVF GTYQLCDLCDNLKTCHTSHGS VMAETAVINHKKRKNSPLIVQS NDLTEAAYSLSRDQKRMLV_F VDQIRKSDGTLQEHDGICGIHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESPFWFIKRSPE*RRGKFPFT QMGQD		l	l		i	1	
LAIDHRGPHISPOARSI, SYNSVE GTYQLCDLCDNILKTCHTSHGS VMAETAVINHKKRKNSPLIVOZ NDLTEAAYSLSRDQKRMLYLF VDQIKKSDGTLQEHDGICEIHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESPFWFIKRSPE*RRGKFPFT QMGQD							
GTYOLCDLONI,KTCHTSHGS			1				
WAMETAVINHKKRKNSPLIVOS NDLTEAAYSLSRDQKRMLYLF VDQIRKSDGTLQEHDGICEHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYFPEEDAGDEK GYESFPWFIKRSPE*RGKFPFT QMGQD 15758 46126 A 15849 1 1773 15750 46128 A 15851 1 726 MPASGNENDLNMPSGTIEIFVR CYVEVERIMSAWTEQPNEHRT MMWSVLLQPCDNLKTCHTSIG SVMAETAVINHKKRKNSPRIVQ SNDLTEAAYSLSRDQKRMLYL FVDQIRKSDGTLQEHDGICEHW AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYFPEEDAGDEK GYESFPWFIKRAHSPSRG/LYSV HINPYLIPFFIGLQNRT/PFRLSE TKEITNPYAMRLYESLCQYRKS 15762 46132 A 15853 1 2121 15763 46131 B 15854 721 2739 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWWAKXYNIPSGLHACTATL EETHILDSGAHVQYFYKGQLH HGYWAETAVSLSRDQKRML HGSWMAETAVINHKKRKNSPRIVQ VMRVPCTDYFDTQKNPDLSHL HGSWMAETAVINHKKRKNSPRIVQ CQWWAKXYNIPSGLHACTATL EETHILDSGAHVQYFYKGQLH HGYWAETAVSLSRDQKRML HGSWMAETAVINHKKRKNSPRIVQ VMRVPCTDYFDTQKNPDLSHL HGSWMAETAVINHKKRKNSPRIVQ CQWWASAVNIPSGLHACTATL EETHILDSGAHVQYFYKGQLH HGYWAETAVSLAEASKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRSAESKDIRQG LGGYGSFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN		1	ı				
NDLTEARYSLSRDOKRMLYLF	1		ŀ				
VDQIRKSIDGTLQEHDGICEIHV				l			
AKYAEJEGLTSAEASKDIRQAL KSFAGKEVVYPYPEEDAGDEK GYESPPWFIKRSP*RGKFPFT QMGQD	1		ļ				NDLTEAAYSLSRDQKRMLYLF
IS758 46126 A 15849 1 1773			l				VDQIRKSDGTLQEHDGICEIHV
GYESFPW/FIKRSPE*RRGKFPFT			1				AKYAEIFGLTSAEASKDIRQAL
New York State							KSFAGKEVVFYRPEEDAGDEK
15758 46126 A 15849 1 1773	ļ	1		ŀ			GYESFPW/FIKRSPE*RRGKFPFT
15759					l		QMGQD
15760 46128	15758	46126	Α	15849	1	1773	
CYVEVERIMSA WTEQPNEHRT MMWSVLLQPCDNLKTCHTSHG SVMAETAVINIKKRKNSPRIVQ SNDLTEAAYSLSRDQKRMLYL FVDQIRKSDGTLQEHDGICEHV AKYAEHFGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESPPWFIKRAHSPSRG/LYSY HINPYLIPFFIGLQNRFTOFRLSE TKEITNPYAMKLYESLCQYRKS TKEITNPYAMKLYESLCQYRKS 15763 46131 B 15853 1 2121 15763 46131 B 15855 1 2529 MQEQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWKAKYNIPSGLHACTATL EFHILDSGAHVQYFYKGQLH VMRVPCTDYFDTQKNPNLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML VLFVPQIRKSDGTLQEHDGICEI HVAKYAEHFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESPPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15856 46134 B 15857 1 411	15759	46127	В	15850	1		
MMWSULLOPCDNILXTCHTSHG SVMAETAVINHKKRKSPRIVQ SNDLTEA4YSLSRDQKRMLYL FVDQIRKSDGTLQEHDGICEIHV AKYABIFGLTSABASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESFPWFIKRAHSPSRG/LYSV HINPYLIPFFIGLQN 15762 46130 A 15853 1 2121 15764 46132 A 15855 1 2739 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWXAKYNIPSGLHACTATL EETHILDSGAHVQVFYKGQLH VMRYPCTDYEDTQKIPNLSHL HGSWMAETAVINHKKRNSPRI VQSNDLTEAAYSLSRDQKBML VLFVQQIRKSDGTLQEHDGICEI HVAKY AEIFGLTSAEASKDIRQ ALSSFAGKEVVFYRPEEDAGD EKGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN	15760	46128	Α	15851	1	726	MPASGNENDLNMPSGTIEIFVR
SVMAETAVINHKRKRNSPRIVQ SNDLTEAAYSLSRDQKRMLYL FVDQIRKSDGTLQEHDGICEHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYFPEEAGDEK GYESPPWFIKRAHSPSRG/LYSV HINPYLIPFFIGLQNRTOFFLSE TKEITNPYAMRLYESLCQYRKS 15762 46130 A 15853 1 2121 15763 46131 B 15854 721 2739							CYVEVERIMSA WTEQPNEHRT
SNDLTEAAYSLSRDOKRMLYL FVDQIRKSDGTLQEIDGICEIHV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GVESPPWFIKRAHSPSRG/LYSV HINPYLIPFFIGLQNRFTQFRLSE TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS 15763 46131	1		ļ				MMWSVLLQPCDNLKTCHTSHG
FVDORKSDGTLQEHDGICEHIV AKYAEIFGLTSAEASKDIRQAL KSFAGKEVVFYRPĒEDAGDEK GYESFPWFIKRAHSPSRGILYSV HINPYLIPFFIGLQNFTOFRLSE TKEITNPYAMRLYESLCQYRKS 15761 46129 B 15852 356 671							SVMAETAVINHKKRKNSPRIVQ
AKYAEJEGLTSAEASKDIRQAL KSFAGKEVVFYRPEEDAGDEK GYESPPWIKKRAHSPSRG/LYSY HINPYLIPFFIGLQNRFTQFRLSE TKEITNPYAMRLYESLCQYRKS HINPYLIPFIGLQNRFTQFRLSE TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS 15762 46130 A 15853 1 2121 15764 46132 A 15855 1 2579 MQEQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWKAKYNIPSGLHACTATL EETHILDSGAHVQVFYKGQLH VMRVPCTDYEDTQKNPALSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML VLFVDQIRKSDGTLQEHGGICE HVAKYAEJFGLTSAEASKDJRQ ALKSFAGKEVVFYRPEEDAGD EKGYESPFWIKKRKHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15765 46134 B 15857 1 411							SNDLTEAAYSLSRDQKRMLYL
RSFAGKEVVFYRPEEDAGDEK GYESFPWFIKRAHSPSRG/LYSY HINPYLIPFFIGLQNRFTQFRLSE TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TKEITNPYAMRLYESLCQYRKS TSF02 46130 A 15853 1 2121 TSF02 46131 B 15854 721 2739 TSF02 AQRTPC-HIRRDGSQRRVCDDCK QQWKAKVNIPSGLHACTATL EETHILDSGAHVQVFYKGQLH VMRYPCTDYEDTQKNPNLSHL HGSVMAETAVINHKKRKNSPR VQSNDCITEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKY AEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESFPWFIKRKAHSPSRGLY SVHINPYLIPFFIGLQN TSF05 46133 B 15856 537 1318 TSF05 46134 B 15857 1 411			l				FVDQIRKSDGTLQEHDGICEIHV
GYESFPWFIKRAHSPSRG/LYSV 15761 46129 B 15852 356 671 15762 46130 A 15853 1 2121 15763 46131 B 15854 721 2739 15764 46132 A 15855 1 5828 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRR VCDDCK QQWWAKYNIPSGLHACTATL EETHILDSGAHVOFFVKQQLH VMRVPCTDYFDTQKNPNLSHL HGSWMAETAVINHKKRKNSPR I VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFGLTSAEASKDIRG ALKSFAGKEVVFYRPEEDAGD EEGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15765 46134 B 15857 1 411			1			ł	AKYAEIFGLTSAEASKDIRQAL
HINPYLIPFFIGLQNRFTQFRLSE 15761 46129 B 15852 356 671 15762 46130 A 15853 1 2121 15763 46131 B 15854 721 2739 15764 46132 A 15855 1 5828 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWKAKYNIPSGLHACTATL EFTHILDSGAHVQYFYKGQLH VMRVPCTDYFDTQKNPNLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML VLFVPQHRSGSGTLQEHGGICEL HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESFPWIRKAKHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318							KSFAGKEVVFYRPEEDAGDEK
TKEITNPYAMRLYESLQYRKS			l				GYESFPWFIKRAHSPSRG/LYSV
15761 46129 B 15852 356 671 15762 46130 A 15853 1 2121 15763 46131 B 15854 721 2739 15764 46132 A 15855 I 5828 MQEQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWWAKYNIPSGLHACTATL EETHILDSGAHVQYFYKGQLH YMRYPCTDYEDTQKNPMLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVYFYRPEEDAGD EKGYESFPWIFKRKHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318		·					HINPYLIPFFIGLQNRFTQFRLSE
15762 46130 A 15853 1 2121				_			TKEITNPYAMRLYESLCQYRKS
15765 46131 B 15854 721 2739 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRRVCDDCK QQWWAKXNIPSGLHACTATL EETHILDSGAHVCPYFKQQLH VMRVPCTDYFDTQKNPNLSHL HGSWMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGVESFPWIFKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15765 46134 B 15857 1 411		46129	В	15852	356		
15764 46132 A 15855 1 5828 MQEQQQASSLIRSGNGRLPGSG AQRTPCHIRRDGSQRYCDDCK QQWKAKYNIPSGLHACTATL EETHILDSGAHVQVFYKGQLH VMRVPCTDYFDTQKNPNLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKKML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESFPWIFKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15765 46134 B 15857 1 411							
AQRTPCHIRRDGSQRRVCDDCK QQWWAKYNIPSGLHACTATL EETHILDSGAHVQVFYKGQLH VMRVPCTDYFDTQKNPNLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAVSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFIGTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGVESFPWFIKRAHSPSGLY SVHINPYLIPFFIGLQN SVHINPYLIPFFIGLQN 13765 46133 B 15856 537 1318 15856 46134 B 15857 1 411 411							
QQWWKAKYNIPSGLHACTATL EETHILDSGAHVQVFYKQCJH VMRVPCTDYFDTQKNPNLSHL HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEJIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN 13765 46133 B 15856 537 1318 13766 46134 B 15857 I 411	15764	46132	Α	15855	1	5828	
EETHILDSGAHVQVFYKGGLH			l			\	
VMRVPCTDYFDTQKNPNLSHL	1		l				
HGSVMAETAVINHKKRKNSPRI VQSNDLIEAAYSLSRDQKRML YLFVDQIRKSDGTQLEDHOGICE HVAKYAEJIGLITSAEASKDIRQ ALKSFAGKEVVFYRFEEDAGD EKGYESFPWIKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 13766 46134 B 15857 1 411		l	l				EETHILDSGAHVQVFYKGQLH
VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIRGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD EKGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN 13765 46133 B 15856 537 1318 13766 46134 B 15857 1 411			l				VMRVPCTDYFDTQKNPNLSHL
YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIRGLTSAEASKDIRQ ALKSFAGKEVVFYPEEDAGD EKGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15766 46134 B 15857 1 411	1		ı	1	l		HGSVMAETAVINHKKRKNSPRI
HVAKYAEIRGLTSAEASKDIRQ ALKSFAGKEVVFYRFEEDAGD EKGYESFPWFIKEAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15766 46134 B 15857 1 411	1		l				VQSNDLTEAAYSLSRDQKRML
HVAKYAEIRGLTSAEASKDIRQ ALKSFAGKEVVFYRFEEDAGD EKGYESFPWFIKEAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1318 15766 46134 B 15857 1 411	1		l				YLFVDQIRKSDGTLQEHDGICEI
ALKSFAGKEVVFYRPEEDAGD EKGYESFPWFIKRAHSPSRGLY SVHINPYLIPFFIGLQN 15765 46133 B 15856 537 1518 15766 46134 B 15857 I 411	1	1	1		l		
EKGYESFPWFIKRAHSPSRGLY	1	1	1		l		
SVHINPYLIPFFIGLQN	1		l		l		
15765 46133 B 15856 537 1318 15766 46134 B 15857 I 411	1				l		
	15765	46133	В	15856	537	1318	1
15767 46135 A 15858 I 1356	15766	46134	В	15857	ı	411	
	15767	46135	Α	15858	1	1356	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		_				
15768	46136	A	15859	1	1215	MPYDRPFTVMAAFPLCPACDK
						EYRDPLDRRFHAQPVACPECGP
						HLEWLKMGKIVAIKGIGGFHLA
					i	CDARNSNAVATLRARKHRPAK
						PLAVMLPVADGLPDAARQLLT
						TPAAPIVLVDKKYVPELCDDIA
						PDLNEVGVMLPANPLQHLLLQ
						ELQCPLVMTSGNLSGKPPAISN
						EQALADLQGIADGFLIHNRDIV
						QRMDDSVVRE\RRNAAPF/AGA
						VLSQHLGDLSDDGIQMQWRER
						YSLMQNIYDFTPQYVVHDAHP
						GYVSSQWAREMNLPTQTVLHH
						HAHAAACLAEHQWPLDGGDVI
						ALTLDGIAQPDERSRSEDVDRA
						HQRGHERDRRRRRSGTSHLTR
İ				l		VDLKEDGKGLKIVRQSLPYGTA
			ĺ			SGTHGLYFCAYCARLHNIEQQL
						LSMFGDTDGKRDAMLRSPNR
15769	46137	В	15860	74	2050	
15770	46138	Α	15861	2	703	LQARFPLSWRNFPITFACYAGL
						FCLSASIIYP/DHLCPVSVPRAFA
						GPRHRRHLLVLHRGCGLRH/TK
						RPGPGPGPARALAIWPPDPGC*
						RCRRPSLPASSLRSSATPTCTST
						QPALEWCVAVYAICFILAAIAIL
				l		LNLGECTNVLPIPFPSFLSALAL
						LSVLLYATALVLWPLYQFDEK
						YGGOPRRSRDVSCSRRHAYYV
1				l		GAWDRRLAEAILTAINLLTYVA
						DLMHSAHLDFSKV
15771	46139	A	15862	I	849	MAQPLAKQWTPVSPRALLVSIS
1		1				SMPPRPAADVTVCRHRNCTLRE
		l	1			LLYGFAYAAAQQLNLHGDVCN
1		l	l	I		DGDG/VEVRLREDPETFLVQLY/
		l	l	I		QHCPPLA\RTGGTMNTQIVPDA
		l	l			ATCPASLSEMNTPGERRYRYPF
	1	l				INCTHYGPSFPIIRAMPYDRPFT
	1	l				VMAAFPLCPAC/DKEYRDPLDR
		l	1			RFHAOP/VACPECGPHLE/WGIG
						GFHLAC/DARNSNAVATLRARK
		1	l			H/RPAKPLAVMLPVADGLPD/A
	1	l	l			
	1	l	l			ARQLLTTPAGAGLWLGGNGYP
		l				RAPDNANSTAMSCASHAHSFG
1.5770	16140	l .	1,6062	l	2688	GGHPRGR
15772	46140	В	15863	1	2688	
15773	46141	A	15864	1	1578	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15774	46142	A	15865	1	877	MQAAHRLRRAYRIKTÖKPPAL ADYHPAYQEMI.SRKQDRVIDQ IDFGIKDTVEILHMGQIQSMGRE FALSLRFSIHGHNVHGNESPEM GPERSGFYDEWSPERIIQYYRP AREQLMNELVQILKNTROHLM TGVSHMIPFVVSGGILLAVSVM LYGKGAVPDAVADPINLKILFD IGVAGLTLMVPFLAAYIGYSIA ERSALAPCAIGAWVGNSFGAGF FGALIDAPFRISEEINPALPEAK VMACLEAVARQDNVDSPQELP PTPLFASRPITRLKSWWAPGGE VESGTHKK
15775	46143	Α	15866	3	423	
15776	46144	A	15867	633	771	NNRSHPALCTNRPPRTRLVFYP LLAPNA*LAALQKGWAYGAGH NA
15777	46145	Α	15868	1	4014	
15778	46146	Α	15869	1	180	SGHPFIPLKDLTQIAIGDVFNNN LILWANREIIHSVP*RY*HDANI QCFVVISSFVRSQQ
15779	46147	A	15870	815	1905	IFQEHINDHLPMPKFGPLAGLR VYPSEMWRAEVIWIENVAWA DTIRVQPNYPQLSRRNLHALSL NIFKDEGREAFLKLMETTDIFIE ASKGPAFARRGITDEVLWEDKP KLVIGHLSGFAHQVSQISTHQG SSNWVSPDQLQCNTQTNSPKQ LESTHCMQCKPNSTTNSRQFQS NPVNIHALNHQSKPHHPLVYAQ LKAKRAHRAKSITASRKQATTF QQVGAPAGGHTHTSSHPQTT KALEQVKHLTTRQAILSKAKPQ TASPIPROTIGPLGRGPHALNPS TTVIGQPANQRIGQHGPGPKP KTPEWPQPKEKVPNRQCTSKTY NLKQOPLNPGKAKT
15780	46148	A	15871	1	361	QTTGISFQGRGGAGPGVPTRTQ/ CLRCHGGRHG/HPSHLCKPNKG DPRKIRLKMSWR*PWFAIGPRD WSSSRPKPTSPRGSCRSFIEASK MSAPVVWSTKTHSSRSMLSFSL MEDFVTALSIL
15781	46149	Α	15872	1	447	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15782	46150	A	15873		1424	MLLEQLSQHKLDMIISDCPIDST QOEGLESVRIGECGYSFWCTINP PPEKPFPACLEERILLIPGRRSM LGRKLLNWFINSQGLONVEILGEF DDAALMKAFGAMHNAIFVAPT LYAYDPYADKTYVEIGRYENV MEEYHAIFAERMIQHPAVORIC UTDYSALFSPALLRRALYHDA TELHPACGKDFRQLYRRIQNQI QRHALWHKIFDVEIPHPILIAR YSANMPHTOCKIAIEWPGKVVE AILRLTNDAPEYLPIGFNIKSK MRSEGMNPYQROFMELALSNQ VLKCGEFTLNRGRKSPYFFNAG LENTGRDLALLARFYAEALVDS GIEFDLLLAPAYKGIPHATTTAV ALAEHHDLDLPYCFNRKEAKD HGEGGRLVGSALQGRYMLVD DVTTAGTAIRESMEIIQANGATL AGVLISLDROEBGR*CDHRRNG DSRYDGDYSGQWRDACWRVD FARSSGTRARRDFGSOWRDACWRVD FARSSGTRARRDFGSOS*A*L QLQSDLYHHPERPDCLPGREAG
15783	46151	В	15874	1	1086	110010000
15784	46152	A	15875	484	729	CTQFRKTCFTCKASNGFCFATS CNLIYTSEIICRGPTG*HNLIASL LQRIDNLRKFIVIPTLKYLTRCR LDDNIMIFYAQGI
15785	46153	A	15876	1717	1971	PANAWAEMWEYSAWQRRMYS MG*TSKVVVHBGATDRW*NTN GCYRIFWIAAPFVPAAVIHGNIL PTHGACVHKGFGRAPAGAAVE
15786	46154	В	15877	I	765	
15787	46155	A	15878	103	823	VPACAGLKKEARSLLASPPRLL TITKLQASCRALFSPPIOSRQTTG ISFQGRGGAOFQVPIRTQCLRC HGGRHG/HPSHLCKFPNKODPRK IRLKMSWR*PWFAIGPROWSS RPRTSPRGSCRSFIEASKMSAP VVWSTKTHSSRSMLSFSLMEM PARMPITSSMSPTELRQGSVMF PARMPITSMSPTELRQGSVMF DFVTALSILLRGTVHEKLRWTF NLYDINKDGYINQEEMMDIVK AIYDMMGKYTYPVLKEDTPRQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
15788	46156	A	15879	1	2926	MVKDASQSQASLGLDAFLPAS
ĺ						GVQYTHERSPDSPLKSQECPSG
						VVNEDTFKQIYAQFFPHGGLMS
				1		ENGTGKKASVFGSTGSGALLLE
		l				EGERRREQPLIEMRKPPHVLIN
						YQAKVGGNSIPSPSLCKPAFPPL
				1		PSWAVDLASARQKEDASRSLQ
						NKEEDFYTSQPGFPALALGCCL
						RVNLLLGHRGISAGSSYRPITST
						PCHLCKLGQEVFAKPLNLQDQ
						CQQNLVMQVLKLVLNCLNFDF
1			ĺ			IGSSADESADDLCTV
15789	46157	A	15880	1	1521	
15790	46158	Α	15881	1	1182	MDKFLDTYTLPRLNQEEVESLN
						RPITGSEIVAIINSLPTKKSPGPD
				l		GFTAEFYQRYKEELVPFLLKLF
				l		QSIEKEGILPNSFYEASIILIPKPG
}				l		RDTTKKENFRPISLMNIDAKILY
}				l		KILAKRIQQHIKKLIHHDQVGFI
ł				l		PGMQGWFNIRKSINVIQHINRA
1				l		KDKNHMIISIDAEKAFDKIQQPF
		1				MLKTLNKLAQNLLKLISNFNKV
						SGYKINVQKSQAFLYTSNRQTE
						SQIMSELPFTIASKRIKYLGIQLT
		1	l			RDVKDLFKENYKPLLKEIKEDT
						NKWKNIPCSWVGRINIVKMAIL
						PKVIYRFNAIPIKLPMTFFTELE
	ŀ					KTTLKFIWNQKRAHIAKSILSQ
				İ		KNKAGGITLPDFKLYYKATVT
						KTAWYCYQNRHIDQWNR\PSEI
						TPHIYNYLIF
15791	46159	Α	15882	1	690	
15792	46160	В	15883	I	458	
15793	46161	Α	15884	1	719	
15794	46162	В	15885	I	3552	
15795	46163	Α	15886	1	321	
15796	46164	A	15887	I	362	
15797	46165	В	15888	1	489	
15798	46166	В	15889	1	1660	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15799	46167	Α	15890	11	1593	MGKKQNRKTGNSKKQSTSPPP
						KERSSSPAMEOSWMENDFVEL
						REEGFRRSNYSELQEDIQTKGK
					Ì	EVENFEKNLEECMTRITNTEKC
						LKELMELKTKARELHEECRSLR
						SRCDQLEERVSAMEDEMNEMK
	ĺ					REGKFREKRIKRNEQSLQEIWD
1		l		ŀ		YVKRPNLHLMGVPESDGENGS
1						KLENTLQDIIQENFPNLARINKI
						DRPLARLIKKKREKNOIDTIKN
						DKGDITTNPTEIQTTIREYYKHL
						YANKLENLEEMDTFLDTYTLPR
		1				LNQEEVESLNRPITGAEIVAIINS
1		1				LPTKKSPGPDGSIAEFYQRYKE
						E/PADKQLQQSLSIQNQCTKITS1
						LIHQQQTNREPNHE*TPIHNCFK
1		l				ENKIPRNPTYKGCEGLLPGELQ
1						TTAQGNKRGYKQTEEHSMLM
		l				GRKNQYRENGHTAQGNF*IQC
l		l				HPHQATNAFLHRIGKNYFKVH
		l				MEPKKSPHRQVNPKPKEQSWR
		l				HHTT*LQTILQGYSNQNSMVLV
1		1				PKQGYRSMEQNRALRNNAAYL
		1				QLSDL*QT*EKHAMGK/EFPI**
15800	46168	Α	15891	1	1177	
15801	46169	Α	15892	1	3663	
15802	46170	Α	15893	1	1995	
15803	46171	Α	15894	1	4191	MVKGSIQQEELTILNTYAAHTG
						APRLIKQVLSDLQRDLDSHTIIM
		l				GDFNTPLSTLDRSTRQKVNKDT
1						QELKSALHQADLTDIYRTLHIIK
		l				STEYTFFSAPHHIYSKIDHILGSK
		l		l		ALLSKCKRTEIITNYLSDHSAIK
		l				LELWIKNLTQNHSTTWELNNLL
						LNDYWVHNEMKAEIKMFFETN
1	1	l				ENKDTTYHNLWDTFKAVCRGK
1	1	l		I	1	FIPLNAHKRKQERSKIDTLTSQL
1		l				KELEKQEQTHSKASRRQEITKIR
		L				AELKEIETQ

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15804	46172	Α	15895	I	1659	MGDFNTPLSTLDRSTRQKVNK
					Ì	DTQELNSALHQADLIDIYRSLH
l						AKSTEYTFFSAPHHTYSKIDHIL
					ĺ	GSKALLSKCKRTEIITNYLSDHS
					ŀ	AIKLELRIKNLTQNRSTTWKLN
ŀ						NLLLNDYWIHDEMKAEIKMFF
					ŀ	ETNENKDTTYQNLWDAFKATA
						SKTNKEKEKNQIDTIKNDKGDI
1						TTDPTEIQTTIREYYKHLYANK
1	l .					LENLEEMDKFLDTYTLPRLNQE
1	1			1		EVESLNRPITGAEIVAIINSLPTK
						KSPGPDGFTAEFYQRYKEELHI
ĺ	1					NRAKDKNHMIISIDAEKAFDKI
						QQPFMLKTLNKLGIDGTYFKIIR
	1					AIYDKPTANIILNGQKLEAFPLK
	1			1		TGTRQGCPLSPLLFNIVLEVLAR
						AIRQEKEIKGIQLGKEEVKLSLF
					1	ADDMTVYLENPIVSAQNLLKLI
						SNFSKVSGYKINVQKSQAFLYT
						NNRQTESQIMSELPFTIASKRIK
	1					YLGIQLTRDVKDLFKENYKPLL
						KEIKEDTNKWKNIPCSWVGRIN
						IMKMAILPKVIYRFNAIPIKLPM
						TFFTELEKTTLKFIWNQK\RARI
15805	46173	A	15896		1740	AKAILSQKNEAGGITLP
15806	46174	A	15897	1	880	
15807	46175	A	15898	í .	2777	MGKKQNRKTGNSKKQSTSPPP
		1	,			KERSSSPAMEQSWMENDFVEL
1						REEGFRRSNYSELQEDIQTKGK
l						EVENFEKNLEECMTRITNTEKC
						LKELMELKTKARELHEECRSLR
1						SRCDQLEERVSAMEDEMNEMK
						REGKFREKRIKRNEQSLQEIWD
						YVKRPNLHLMGVPESDGENGS
	l					KLENTLODIIQENFPNLASTLSD
						HSGIKLEINSKRNPONHANTWK
				1		LSNLLLNAHWVNNEIKMKVKN
						FFELNDNSDTIYQNFSD
15808	46176	Α	15899	326	3096	QTERTSTPKTHLYVTIIKDQK*I
	1					KPQRRGKNRGEKLETLKSRAP
						VLLQRNAVPQQQRNKAGRRM
				1		TLTS*EKKSSDNQTTPSYRRKF
l	1	1	1		1	KPMAKKLKTL/RKKLDECITRIT
						NAEKCLMELMELKAKARELRE
	1	1	1			ECKSLRSQCNQLEER\INKIDRP
	1		1			LARLIKRKREKNQIDAIKIDKGD
						ITTOPTEIQTTIREYYKHLYANK
				l		LENLEEMDKFLDTYTLPRLNQE
	1	1	1		l	EVESLNRPITGAEIVAIINSLPTK
				1		KSPGPDGFTA
		_				

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hed	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, *=possible nucleotide insertion)
15809	46177	Α	15900	li .	2047	MENDFEELREEGFRRSNYSELR
						EDIQTKGKEVENFEKNLEECITR
i		l				ITNTQKCLKELMELKTKARELR
l					1	EECRSLRSRCDQLEERVSAMED
l						EMNEMKREGKFREKRIQRNEQ
i						SLQEIWDYVKRPNLRLIGVPET
l						LHPRDEAHLKNLIDAIKNDKGD
l						ITTDPTEIQTTIREYYKHLYANK
İ	1	l				LENLEETDKFLDTYTLPRLNQE
l						EVESLNRPITGSEIVAIINSLPTK
l	1	ł				KSPGPDGFTAEFYQRYKEEL/PD
i	1					KQLQQSLRIQNQCTKITSILIHQ
i	İ	ĺ		1		QQTNREPNHE*TPIHNCFKENKI
	i					PRNPTYKGCEGPLQGELQTTAQ
ĺ	1					GNKRGYKQMEEHSMLMGRKN
ĺ	1					QYHENGHTAQGNLQIQCHPH*
	1	l				ATNAFLHRIGKNYFKVHMEPK
	1	ł		1		KSPHRQVNPKPKEQSWRHHTT
	1			ļ		*LQTILQGYSNQNSMVLVPKQR
	i					YRSMEQNRALRNNAAYLQLSD
	i	1				L*QT*EKQEMGKGFPI**MVLG
	İ	l		l		KLASHM*KAETGSLPYTLYKN
	l	1				QFKMD*RLKR*T*NHKNPRRKP
	1					RHYHSGHRHGQGLHV*NTKSN
						GNKSQN*QMGSN*TKELLHSK
i						KKKNY*QTEQATCKMGENFHN
						LLI*QRANIQNLQ*TQTNLQEKN
	1		1	ŀ		KQPHQKVGKGHEQTLLKRRHL
				l		CSQK\NHEKMVTITGHQRNAN
						QNHNEIPSHAS*NGNH*KVRKQ
				l		VLERMCRNRNTFTLLVGL*TSS
15810	46178	Α	15901	1	3285	
15811	46179	A	15902	1	3681	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclentide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codnn, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15812	46180	Α	15903	1	1854	MKAEIKTFFETNENKDTTYONL
		1		ŀ		WDAFKAVCRGKFTALNAHKR
		l				KQERSKIDTLTSQLKELEKQEQ
		l				THSKASRRQEITKIRAELKEIET
		l				QKALQKINESRSWCFEKIHKID
		l				RPLARLIKKKREKNQIDAIKND
		l				KGDITTDPTEIQTTIREYYKHLY
		ı				ANKLENLEEMDKFLDTYTLPRL
						NQEEVESLNRPITGSEIVAIINSL
		1				PTKNSPGPDGFTAEFYQRYKEE/
		l				RENKIPRNPTYKGCEGPLQGEL
		ı				QTTAQGNKRGYKQMEEHSML
İ		1				MGRKNQYRENGHTAQGNLQIQ
		1		ŀ		CHPHQATNDFLHRIGKNYFKV
		1		1		HMEPKKSPHRQVNPKPKEQSW
		1		l		RHHTT*LQTILQGYSNQNSMVL
1		1				VPKQRYRSTEQNRALRNNATY
		1		ļ		LQLSDL*QT*EKQAMEKGFPI**
		1				MVLGKLASHM*KAETGSLPYT
				ļ		LYKNQFKMD*RLKR*T*NHKN
		1				PRRKPRHYHSGHRHGQGLHV*
				l		NTKSNGNKSQN*QMGSN*TKE
		1				LLHSKRNYHQSEQATYKMGEN
		1				FRNLLI*QRANIQNLQ*TQTNLQ
		1		l		EKNKPPHQKVDEGHEQTLLKR
		1				RHLCSQKTHEKMLIITGHQRNA
		1				NQNHNEIPSHTS*NGNH*KVRK
		1		i		QQVLERMWRNRNTFTLLVGL*
15010	46101	<u></u>	15904		2718	TSSTIVEVSVV
15813	46181	B A	15904	1896	5031	TWKGTTSTSRCKIMPKYRSTRO
15814	40182	l ^A	13903	1890	3031	KVNKDTOELNSALHOADLIDIY
		1				RTLHTKSTEYTFFSAPHHTYSK
		1				TDHIVGSKALLSKCKRTEIITNY
		l				LSDHSAIKLELRIKNPTOSRSTT
		l		ŀ		WKLNNLLLNDYWVHNEMKAE
		1	1			IKMFFETNENKDTTYONLWDT
		l				FKAVCRGKFIALYAHKRKQERS
		l	i	İ		KIDTLTSQLKELEKQAQTHSKA
1		l	1	l		SRROEITKIRAELKEIETOKTL\Q
			ĺ	ĺ		KINE\SRSWFF\ERINKIDRPLAR
1		l	l			LIKKKREKNO
15815	46183	A	15906	4	133	LIGUIGERING
13013	C0101	I**	13300	<u> </u>	1.50	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15816	46184	Α	15907	95	1875	PFFGRKNYRLPSKNMPRRCRSS
						PLPCSMKMTI*CVLTSTATITRL
						PAASAQW*LDGRTLVTKNSFR
				1		YLHTLHTMGPAPEPNLTILWSE
						ELPIAFKKYAAQVSIVTSSLQYE
				l		NDDLMRTDFNSDDYAIACCVS
	į	1		l		PMVIGKQMQFFGARANLAKTL
				l		LYAINGGVDEKLKIQVGPKTAP
		1		l		LMDDVLDYDKVMDSLDHFMD
		1				WLAVQYISALNIIHYMHDKYSY
		l		l		EASLMALHDRDVYRTMACGIA
		1			1	GLSVATDSLSAIKYARVKPIRD
		1				ENGLAKDIQGTTEPQFRNSHSR
1		1				TSGTVTLYLPVSTFVGTGFFIRV
		1		1		DPPKKPLKIRKIEFVEIYEPFSEG
						QAKAQADRCLSCGNPYCEWKC
		1				PVHNYIPNWLKLANEGRIFEAA
		1	İ	ŀ		ELSHQTNTLPEVCGRVCPQDRL
		l				CEGSCTLNDEFGAVTIGNIERYI
		l	İ			NDKAFEMGWRPDMSGVKQTG
		l	1	l		KKVAIIGAGPAGLACADVLTRN
l		1				GVKAVVFDRHPEIGGLLTFGIP
						AFKLEKEVMTRRREIFTGMGIE
						FKLNTEVGRDPRASSFYEEKQL
		ŀ				IARKALPFIDNKTTMAADSSST
						VMELLKLLQDRSGLTLLTNSAE
1						AIHVLAQSEIKVVSTGGELNKN
	1					TLSLQGRITKEIIRRYHVDIMWL
	1					PDELRVEKGFSPDTLKLLEAKG
						QKVALKEAMGSTQSIMVGPDE
	46105	_	1.5000	071	0110	RRTGFSYLRYQNLMIRVKF
15817	46185 46186	B A	15908	271 597	2118	
15819	46187	A	15910	573	1066	IRAQNGSPWLSFLVDSVCVLKA
13019	40107	1	13710	373	1000	CVDAVMRAGICSVS\YWQYQS
						GVGANSSALSGFSOGAIMMLES
i						IKAEPGLASRVIAFNGRYSSLPE
1		1			i	TASTATTIHLIHGGEDPVIDLAH
1		1				AVAAQEALISAGGDVTLDIVED
1		1				LGHAIDNRSMQFALDHLRYTIP
						KHYFDEALSG
15820	46188	A	15911	1	2652	
15821	46189	В	15912	1	1494	
	1	드	1	L		l

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15822	46190	A	15913	li .	1733	MGVGSKGTIFGSTTTQHGSFIPY
			i			LTGYLTYTHGHSPANPKPAVG
1						GOILTAVLPAHDGYVSRIMTM
1						NTOLTTPLTTATHTAROTASLIS
						VIGRQLRCFRHSSQPSVLTHVR
						DDSTQYAQLDMANGQPTSHGY
				1		EHWAOLMYATDRRYGRHMSD
						AARQGCSDTGIVKFSHPEPEMS
				1		VRKFTSARSGCVRSGAWNVTV
						SAQTRVSQALTTLLRRVHKSPM
1						NALTVPAVMAEAPSOAHGRRO
1						TRRLSLHQTLRLLLTTPDWVGV
						IRRCVPVVWGCLEHLDTAGFPL
				1		PNSHTMPLPIGWNDTRKRLHHS
						AHNGQHDVAQGRRVLGRKVG
1						LTHPKVQQQLGVDGLSLFCVY
	1					LVWYIGLHLRRPFRLAYTICPSL
1					i	VYGADLRVIVRLFSDSHHFMAF
1				ł		GNPQPKHLDCQEYALFLLNSM
						PDNTAEHYRNKIAIYLHCLPED
			ĺ	ŀ		DRIKAINEIRMAIHQVSPFREEP
				ł		VDCVLWVKYSQLMPNDYNPN
				1		NVAPPEKKLLQKSIEIDGFTQPI
				į .		VVTHTDKNAMEIVDGFHRHEI
				ŀ		GKG/SHASLKLRLRGYLPVTCL
						EGTRNQRIAATISHNRARGRHH
					l	ITAMSEVTPELRLLGW
15823	46191	В	15914	I	273	
15824	46192	В	15915	169	292	
15825	46193	A	15916	2	844	KGGVYKTSVSVHLAQDLALKG
				ł		LRCLLWKGT\PPQGTPSMVSRE
				l		GDPALHN*CKTQLLCLSYLGEK
						DDVTYAIKPTCWPGLDIIPFCLA
						LDRIETEL\MGKFDEGKLPTDPH
	'			1		LMLRLAIETVAHDYDVIVI\DSA
				1		HNRG\IGTINVVCAADVLIVPTP
				ł		VE\MFDYTSALQFFDMLRDLLK
				1	1	NA\DLKGFEPDVRIFVTKYSNSN
		1		l		GSOFP\WMEEPIPECLPKHGSKK
				l		NNVRNANEVGKGQIRMRT\VFE
						OAIDORSSTGAWRNALSIWEPV
1		l	1	l		CN/EIFDRLIKPRWEIR
	L			L	L	CITIES DICIKI KWEIK

15826 46194 A 15917 I 681 MSIGGSGNSLCTRPKQQQGGT LKQYEVSVPGTLRCLACPQAC LKREHWHFAPKSNYDSCRYIE EYFAYEASFMSVQTELVIKSS EPLASLEMHIKKNPVSIPHTVW YADDIRRGEREAADVLGLTLY LMLRAGEAAPDVLGTLY LARAWGIDDTLAGESDKPHF HWLVLCGHGNNGGDGYVVAI LARAWGIDDTLAGESDKPHF LARAWGIDTLAGESDKPHF LARAWGIDDTLAGESDKPHF LARAWGIDTLAG	SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
15826 46194 A 15917 I 681	NO:		hod			codon for last amino acid	
IS826 46194 A IS917 I 681 MSIGGSGNSLCTRPKQQQGGT LKQYEVSYPGTLRCLACPQAC LKKEHWHFAPKSNYDSCRYIE EYFAYEASHMSVQTELVIKSE EPLASLEMHIKKNPVSIPHTVW YADDIRRGEREAADVLGLTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LMLRAGEAAPDVLGTLY, LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLLY LAMPSEHLY L		sequence		09/540,217		of peptide sequence	deletion, \"possible nucleotide insertion)
LKQYEVSVPGTLRCLACPQAC LEKREHWHFAPKSNYDSCRYIE EYFAYEASFMSVQTELVIKSI EDLASLEMHKKNPVSIPHTVW YADDIRRGGERADVLGLTLY LMLRAGEAAPQVCRSAYPDTR HWLVLCGHGNNGGDGYVVAF LAKAVGIDDTLAQESDKPJPE EAPLAHEAWVNAGGEHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGQFSOFIQKL VIFLDPLFIRKQQGVTPTAFAM HLHEVISQGLESILGALDIEGSS DKQRITITATTPSVGALVLPVIN RAIKTHYPQLLLRIPPISDARM LSQFGTDLIDNMFCTNRTVQ-H HGLFAANMVLICREGNPLLSLI DDREAIANAAHVLLIPEQMPS GLRQRVQEMPFGRQINFTSYNI LTIAALVANDDMALIPSRLYNI PSRCWPLEKLPHPSLNEQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISELMSLGIGGVVAVGVIVASV VLFLRVLSMSWLKEVIGTEKA* IAMCHLRALPGOPSFDAQLGM NWVIDKAWDDLMALQNGGVI AMFINERSLPYLTKGLVPALV ARELLVRGMYVRGRMVAFLI CCVALAISAMYELIEWWAALA MGQGADPHLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNESLPYLTKGLVPALV MGQADPHLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNESLPYLTKGLGVPALV MGRINGFGGDGWDTGGGGVDVVMGGGVDV VMFSNESLPYLTKGLGVPALV MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFRIEFT MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFRIEFT GAYANDFGVWDTNVGETIRHET GAYANDFGVWDTNVGETIRHET GAYANDFGVWDTNVGETIRHET					sequence		
LEKEHWHIFAPKSNYDSCRVIE EYFAYEASFMSVQLTELVIKSL EDLASLEMIHKKNPVSIPHTYW YADDIRRGEREAADVLGITLY LMLRAGEAAFQVCRSAYPDTS HWLVLCGHGNNGGDGYVVAI LAKAVGIDDTLLAQESDKPLP EAPLAHEAWYNAGGEHALNI GWPESEHLIV VKCSEKCLFLPLGQPSQFIQKL NIFELDPLFIRRQQVTPTAFAM HILIEVISQGLESILGALDIEGSY DKQRTITIATTPSVGALVLPIVI RAIKTHYPQLLLRNPPISDAEN LSQFQTDLIIDNMFCTNRTVQH HGLFAANMVLLCREGONPLLSL DDREAIANAAHVLLPEQQNF GLRQVQEMFPGQNIFTSYN LTTIAALVANDMLAIPSELYNI PSRCWPLEKLPHPSLNEEQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGO SISELMSLGIIGVVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHERALFOOPSFDAQLGM NWVIDKAWDDLMALONGGVI AVMFSNEESLPYLTKGLVPALA MGQADDFLGTQGQWDTTOS GTYLRAAWRINDGDIPRSLSLE PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALONGGVI VMFSNESLPSTLTKUPFETTAA MARIIGGLMSDIRIPFGVNVLW DDVASFDELAMATGAKFIEIFT GAYASDFGVWDTNVGETIRRIET GAYASDFGVWDTNVGETIRRIEFT	15826	46194	Α	15917	1	681	MSIGGSGNSLCTRPKQQQGGTG
EFFAYEASFASFOUCTELVIKSL EDLASLFMIHKKNPVSIPHTVW YADDIRRGEREADVLGLTLY LMLRAGEAFQVCRSAYPDTF HWLVLCGHGNNGGGVVVAA LAKAVGIDTLLAQESDKPLPE EAPLAHEAWVNAGGEHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGGFSQFIQKL VIVIELDPLFIRKQGVTPTAFAM HLHEYISQGLESILGALDIEGSV DKQRITITATTPSVGALVLPVIV RAIKTHYPQLLLRNPPISDAEN LSQFGTDLIIDNMFCTNRTVQ-H HGLFAANMVLLLPEQRFS GLRQRVQEMFPGRQINFTSYNI LTIAALVANSDMLAIIPSRLYNI PSRCWPLEKLPHPSLNEEQIDF HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGSAQISALSVGNMFOV SISELMSLGIIGGVVAVGIVASS VLFLRVLSMSWLKEVIGTEKA LAMCHERALFGOPSFDAQLGM NWVIDKAWDDLMALONGGVI AVMFSNEFSLPVLTKGLVPALV AREILVRGMYVRGRRMVAFLL CCVALAISAMYELIEWWAALA MGQGADPLGTGGQDWDTOS GYVLRAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALFGDFSFDAQLGMN WVIDKAWDDLMALONGGVU VMFSNEFSLPLTKTGLOFANL MGGADDFLGTGGDQWDTOS GYVLRAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALFGDFSFDAQLGMN WVIDKAWDDLMALONGGVU VMFSNEFSLPLTKTGLOFANL MGGADDFLGTGGDQWDTOS GYVLRAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALFGDGSFDAQLGMN WVIDKAWDDLMALONGGVU VMFSNEFSLPLTKTGROFANL MRIIGGLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIRIFT GAYYASDFOVWDTNVGETIRHEIT GAYASDFOVWDTNVGETIRHEIFT GAYASDFOVWDTNVGETIRHEIFT							LKOYEVSVPGTLRCLACPOACT
EDLASLEMIHKKNPVSIPHTVW YADDIRRGEREAADVLGLTLY LMLRAGEARQVCRSAYPDIR HWLVLCGHGNNGGDGYVVAI LAKAYGIDDTLAQESNCPLP EAPLAHEAWVNAGGEHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGQFSQFIQKL NVIFLDPLFIRRGQGVTPTAFAA HLIEVISQGLSELIGADIDEGSS DKQRTITIATTPSVGALVLPVIN RAIKTHYPQLLLRIAPPISDABAN LSQFQTDLIIDNMFCTNRTVQH HGLFAANNVLICREGNPLLSLI DDREAIANAAHVLLPFEQMPF GLRQEVQEMPFGQINFTSYNI LITIAALYANSDMLAIIPSRLVNI PSRCWPLEKLPHPSLNEQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSPAGESAQISALSVGNMFGV SISSELMSLGIIGVVAVGVIAVS VLFLRVLSMSWLKEVIGTEKA* LAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGYV AMFSNETSLPYLTKGLVPAL AREILVRGMYVRGRMVAFLI CCVALAISAMYELIEWAALA MGQGADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKGLVPALV AREILVRGMYVRGRMVAFLI CCVALAISAMYELIEWAALA MGGQADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKGLGVFAL MGGADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKQRGTMAAT MGRIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFRIEIT GAYASDFGVWDTNVGETRIREIT GAYASDFGVWDTNVGETRIREIT GAYASDFGVWDTNVGETRIREIT							LRKEHWIHFAPKSNYDSCRYIE
EDLASLEMIHKKNPVSIPHTVW YADDIRRGEREAADVLGLTLY LMLRAGEARQVCRSAYPDIR HWLVLCGHGNNGGDGYVVAI LAKAYGIDDTLAQESNCPLP EAPLAHEAWVNAGGEHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGQFSQFIQKL NVIFLDPLFIRRGQGVTPTAFAA HLIEVISQGLSELIGADIDEGSS DKQRTITIATTPSVGALVLPVIN RAIKTHYPQLLLRIAPPISDABAN LSQFQTDLIIDNMFCTNRTVQH HGLFAANNVLICREGNPLLSLI DDREAIANAAHVLLPFEQMPF GLRQEVQEMPFGQINFTSYNI LITIAALYANSDMLAIIPSRLVNI PSRCWPLEKLPHPSLNEQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSPAGESAQISALSVGNMFGV SISSELMSLGIIGVVAVGVIAVS VLFLRVLSMSWLKEVIGTEKA* LAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGYV AMFSNETSLPYLTKGLVPAL AREILVRGMYVRGRMVAFLI CCVALAISAMYELIEWAALA MGQGADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKGLVPALV AREILVRGMYVRGRMVAFLI CCVALAISAMYELIEWAALA MGGQADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKGLGVFAL MGGADDFLGTGQDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGYD VMFSNETSLPYLTKQRGTMAAT MGRIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFRIEIT GAYASDFGVWDTNVGETRIREIT GAYASDFGVWDTNVGETRIREIT GAYASDFGVWDTNVGETRIREIT				i			EYFAYEASFMSVOLTELVIKSL
LIMERGGEAFGVCRSAYPOTE HWLVLCGHGNNGGDGYVVAI LAKAVGIDDTLLAQESDKPLPE EAPLAHEAWVNAGGEIHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGOFSOFIOKL VIVELDPLFIRKOGGVTPTAFAM HUTEVISQGLESILGALDIEGSY DKQRITITATTPSVGALVLPVIN RAIKTHYPQLLLRIPPISDARM LQQFOTDLIDNMECTNRTVQH HGLFAANMYLICREGNPLLSLI DDREAIANAAHVLLLPEGOPS' GLRQRVQEMPFGRQINFTSYNI LTIAALVANDMALIPSRLYNI PSRCWPLEKLPHPSLNECQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISSELMSLGIGGVVAVGVIAVSV VLFLRVLSMSWLKEVIGTEKA' IAMCHEALFOOPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPYLTKGLVFALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGQGADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVFALV MGQADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLGVFALV MGCADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALFOOPSFDAQLGM WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLGVFALV MGRINGFONVILW DPVASFDLAMATGAKFRIEFT GAYASDFGVWNLW DPVASFDLAMATGAKFRIEFT GAYASDFGVWNLW	l	1					EDLASLFMIHKKNPVSIPHTVW
LIMERGGEAFGVCRSAYPOTE HWLVLCGHGNNGGDGYVVAI LAKAVGIDDTLLAQESDKPLPE EAPLAHEAWVNAGGEIHALNI GWPESEHLIV 15827 46195 A 15918 3 774 VKCSEKCLFLPLGOFSOFIOKL VIVELDPLFIRKOGGVTPTAFAM HUTEVISQGLESILGALDIEGSY DKQRITITATTPSVGALVLPVIN RAIKTHYPQLLLRIPPISDARM LQQFOTDLIDNMECTNRTVQH HGLFAANMYLICREGNPLLSLI DDREAIANAAHVLLLPEGOPS' GLRQRVQEMPFGRQINFTSYNI LTIAALVANDMALIPSRLYNI PSRCWPLEKLPHPSLNECQIDF: HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISSELMSLGIGGVVAVGVIAVSV VLFLRVLSMSWLKEVIGTEKA' IAMCHEALFOOPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPYLTKGLVFALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGQGADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVFALV MGQADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLGVFALV MGCADDFLGTGOGDWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALFOOPSFDAQLGM WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLGVFALV MGRINGFONVILW DPVASFDLAMATGAKFRIEFT GAYASDFGVWNLW DPVASFDLAMATGAKFRIEFT GAYASDFGVWNLW							YADDIRRGEREAADVLGLTLYE
HWLVLCGHGNNGGDGVVVAI LAKAVGIDDTLLAQESDKPLPE EAPLAHEAWVNAGGEIHALN EAPLAHEAWVNAGGEIHALN GWPESEHLIV VKCSEKCLEPLGOPSOFIOKL VIFLDPLFIRKGQGVTPTAFAM HILTEVISQGLESILGALDIBGSS DKORTITIATTPSVGALVLPIVI RAIKTHYPQLLLRNPPISDAEN LSQFOTDLIDNMFCTNRTVOH HGLFAANMVLICREGNPLLSLI DDREAIANAAHVLLLPEEQNPS GLRQRVQEMPEQROIPTISVN LTIAALVANDMLAIPSRLYNI PSRCWPLEKLPHPSLNEEQIDS HYNKFSLR HYNKFSLR MITVLLIATQFAPYFTEMALKC GFSFAGESAGISALSVGNMFG SISELMSLGIIGVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHERALPGDPSFDAQLGM NWIDKAWDDLMALONGGVI AVMSNEESLPYLTKGLVPAL CCVALAISAMYELIEWWAALA MGQADPLGTQGQWDTTOG GYYLRAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNESLPLYTKVRFETTAA MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNESLPLYTKVRFETTAA MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNESLPLYTKVRFETTAA MARIIGGLMSDIRIPFGVNVLW DDVASFDLAMATGAKFIRIFT GAYASDFGVWDTNVGETIRHE GAYASDFGVWDTNGTMRHE GAYASDFGVWD	1						
EAPLAHEAWYNAGGEIHALNI GWPPESEHLIV							HWLVLCGHGNNGGDGYVVAR
EAPLAHEAWYNAGGEIHALNI GWPPESEHLIV			ŀ				LAKAVGIDDTLLAQESDKPLPE
15827 46195 A 15918 3 774			ŀ				EAPLAHEAWVNAGGEIHALNI
IVIELDPLFIRKGQGVTFTAFAM HLITEVISQLESILGALDIEGSY DKQRTITIATTPSVGALVLPVIY RAIKTHYPQLLIRNPPISDAEN LSQFOTDLIDNMFCTNRTVQH HGLFAANMVLICREGNPLLSLI DDREAIANAAHVLLLPEEQNF: GLRQRVQEMFPGQINFTSYNI LTIAALVANSDMLAIIPSRLYNI PSRCWPLEKLPHPSLNEEQIDF: HHYNKFSLR HTYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISELMSLGIIGVVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEESLPYLTKGLVPALI CCVALAISAMYELIEWWAALA MGQGADPLGTQGQWDTTOG GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEESLPYLTKVRGETTAAL MARIIGGLMSDIRIPFGVNVLW DVASFDLAMATGAKFRIEFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT							
IS828 46196 A IS919 I 741 MITVLLIATQFAPYTEMALKC GFSFAGESAQISALSYONMEN 15828 46196 A IS919 I 741 MITVLIATQFAPYTEMALKC GFSFAGESAQISALSYONMEN IS828 46196 A IS919 I 741 MITVLIATQFAPYTEMALKC GFSFAGESAQISALSYONMEN SISELMSLGIIGVVAVGVASS VLFLRVLSMSWLKEVIGTEKA IAMCHLRALFGDPSFDAQLGM NWVIDKAWDDLMALONGGVI AVMFSNESSLPYLTGLUPALI CCVALAISAMYELIEWWAALA MGQGADPLGTGGQDWDTOG GYYLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A IS920 I 678 MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNESSLPYLTKUFVATA MARIIGGLMSDIRTPGVNVLW DPVASFDLAMATGAKFRIEFT GAYASDFGVWDTNVGETIRRIEFT GAYASDFGVWDTNVGETIRRIEFT GAYASDFGVWDTNVGETIRRIEFT	15827	46195	A	15918	3	774	VKCSEKCLFLPLGQFSQFIQKLR
DKQRTITIATTPSVGALVLPVIN RAIKTHYPQLLLRNPPISDAEN LSQPQTDLIIDNMFCTNRTVQH HGLFAANMYLICREGNPLLSLI DDREAIANAAHVLLPEEQNPF GLRQRVQEMPFGRQINFTSYNI LITIAALYANDBMALIPSRLYNI PSRCWPLEKLPHPSLNEEQIDP: HHYNKFSLR 15828 46196 A 15919 I 741 MITVULIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISSELMSLGIIGVVAVGVIAVSI VFLRVLSMSWLKEVIGTEKA* LAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGVVI AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLE 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVPALV ARGILVRGMYVRGRKMVAFLI CVALAISAMYELIEWWAALA MGQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLE AMGULKALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVPALV ARGILVRGMYVRLTKVRFETTA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKRRIEIT GAYASDFGVWDTNVGETIRKEIT GAYASDFGVWDTNVGETIRKEIT GAYASDFGVWDTNVGETIRKEIT			1				VIFLDPLFIRKGOGVTPTAFAM
DKQRTITIATTPSVGALVLPVIN RAIKTHYPQLLLRNPPISDAEN LSQPQTDLIIDNMFCTNRTVQH HGLFAANMYLICREGNPLLSLI DDREAIANAAHVLLPEEQNPF GLRQRVQEMPFGRQINFTSYNI LITIAALYANDBMALIPSRLYNI PSRCWPLEKLPHPSLNEEQIDP: HHYNKFSLR 15828 46196 A 15919 I 741 MITVULIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISSELMSLGIIGVVAVGVIAVSI VFLRVLSMSWLKEVIGTEKA* LAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGVVI AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLE 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVPALV ARGILVRGMYVRGRKMVAFLI CVALAISAMYELIEWWAALA MGQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLE AMGULKALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKGLVPALV ARGILVRGMYVRLTKVRFETTA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKRRIEIT GAYASDFGVWDTNVGETIRKEIT GAYASDFGVWDTNVGETIRKEIT GAYASDFGVWDTNVGETIRKEIT							HLHEYISOGLESILGALDIEGSY
RAIKTHYPQLLERNPPISDAEN LSQFQTDLIIDNMFCTNRTVQH HGLFAANMVLLCREGOPLUSLI DDREAIANAAHVLLPEEQMF GLRQRVQEMFPGQINFTSYNI LTIAALVANSDMLAIIPSRLYNI PSRCWPLEKLPIPPSLNEEQIDF: HHYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVONMFGC GFSFAGESAQISALSVONMFGC SISELMSLGIIGVVVAVGIVASV VUFLRVLSMSWLKEVIGTEKA IAMCHERALPGDFSFDAQLGM NWUIDKAWDDLMALONGGVI AVMFSNEESLPYLTKGUPAJA AREILVRGMYVRGRKMVAFLV CCVALAISAMYELIEWWAALA MGQGADPLGTQGQWDTTOS GYVLRAAWRINDGDIPRSLSLE PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNESLPYLTKUFVETAA MARIIGGLMSDIRIPFGVNVLW DDVASFDLAMATGAKFREIFT GAYASDFGVWDTNVGETIRRIEFT GAYASDFGVWDTNVGETIRRIEFT			l				DKORTITIATTPSVGALVLPVIY
HGLFÄANMYLICREGRPLLSLI DDREAINAAHVLLIPEEQNF: GLRQRVQEMFPGQINFTSYNI LTTIAALVANSDMLAIIPSRLYNI PSRCWPLEKLPHPSLNEQIDF: HYNKFSLR HYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSVGNMFGV SISELMSLGIIGVVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHLRALFGDPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNESLPYLTKGLVPALI CCVALAISAMYELIEWWAALA MGQADPLGTQGQWDTTGG GYYLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNESLPYLTKVRGETTAAL MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT			1				RAIKTHYPOLLLRNPPISDAENO
DDREAIANAAHULLIPEGONFI GLRQRVQEMFPGRQINFTSYNI LTIAALVANSDMLAHIPSRLYNI LTIAALVANSDMLAHIPSRLYNI PSRCWPLEKLPHPSLNEEQIDFI HHYNKFSLR 15828 46196 A 15919 I 741 MITVLLIATQFAPYFTEMALKC GFSFAGESAQISALSYGNMFGV SISELMSLGHIGGVVAVGIVASS VLFLRVLSMSWLKEVIGTEKA IAMCHERALFGOPSFDAQLGM NWVIDKAWDDLMALONGGVI AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGQADDFLGTGGODWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNEFSLPYLTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DDVANSTELABLYTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DDVANSTELABLYTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DDVANSDFSLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT	l		l				LSQFQTDLIIDNMFCTNRTVQH
GLRQRVQEMFPGRQINFTSYNI LTTAALVANSDMLAIIPSRLYNI			ĺ				HGLFAANMVLICREGNPLLSLE
15828 46196 A 15919 I 741 MITVLIATQFAPYFEMALKG GFSFAGESAQISALSVGNMGV SISELMSLGIIGVVVAVGIVASV VUFLRVLSMSWLKEVIGTEKA IAMCHERALFODFSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEESLPYLTKGLVPAL) AREILVRGMYVRGRKMVAFLA CCVALAISAMYELIEWWAALA MGQGADPLGTQGQWDTOS GYVLRA WEINDGDIPRSLSLE PATALWLDNII 15829 46197 A 15920 I 678 MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNESLPYLTKUFVALA ARRIGQLMSDIRTFGVVVLW DVVMFSNESLPYLTKVRFETTAA MARIIGGLMSDIRTPGVVVLW DDVASFDLAMATGAKFREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT	1		1				DDREAIANAAHVLLLPEEQNFS
FSRCWPLEKLPHPSLNEEQIDF: IHYNKFSLR	1		1				GLRORVQEMFPGRQINFTSYNI
IHYNKFSLR	1		l				LTIAALVANSDMLAIIPSRLYNL
15828 46196 A 15919 I 74I MITVLILATOFAPYFTEMALKC GFSFAGESAQISALSYGNMFOV SISIELMSLGIIGGVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHERALFODFSFDAQLGM NWVIDKAWDDLMALONGGVI AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGGADPLGTGGDQWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI MCHLRALFGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNEFSLPYLTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT	1		i		}		FSRCWPLEKLPHPSLNEEQIDFS
GFSFAGESAQISALSVGNMFGV SISELMSLGIIGVVVAVGVGVASV VLFLRVLSMSWLKEVIGTEKA* IAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPVLTKGLVPALJ AREILVRGMYVRGRKMYAFLI CCVALAISAMYELIEWWALAL MQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD/ VMFSNEFSLPYLTKVRFETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT			1				IHYNKFSLR
SISELMSLGIIGVVAVGIVASV VLFLRVLSMSWLKEVIGTEKA IAMCHLRALPGDPSPDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPYLTIGLUPAL) AREILVRGMYVRGRKMVAFLV CCVALAISAMYELIEWWAALA MGGADPLGTGQDQWDTOS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKVRFETTAA MARIIGGLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT	15828	46196	Α	15919	I	741	MITVLLIATQFAPYFTEMALKG
VERRVLSMSWLKEVIGTEKA* IAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGYU AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGGOADPLGTGODQWDTQS GYVLRAAWRINDGDIPRSLSLY PATALWLDNRI MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD VMFSNEFSLPYLTKVRFETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKRIEIT GAYASDFGVWDTNVGETIRHET GAYASDFGVWDTNVGETIRHET			1		Į	1	GFSFAGESAQISALSVGNMFGW
IAMCHLRALPGDPSFDAQLGM NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPYLTKGLVPAL) ARBILVRGMYVRGRKMYAFLI CCVALAISAMYELIEWWAALA MGGADDFLGTQGDQWDTGS GYYLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD/ VMFSNEFSLPYLTKVRFETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT			1				SISELMSLGIIGVVVAVGIVASV
NWVIDKAWDDLMALQNGGVI AVMFSNEFSLPYLTKGLVPALV AREILVRGMYVRGRKMVAFLI CCVALAISAMYELIEWWAALA MGGADDFLGTGGDQWDTGS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD. VMFSNEFSLPYLTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT GAYASDFGVWDTNVGETIRHEIFT			1		Ì		VLFLRVLSMSWLKEVIGTEKAV
AVMFSNEFSLPYLTKGLVPALI AREILVRGMYVGRKWAFLI CCVALAISAMYELIEWWAALA MGGGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALQNGGVD, VMFSNEFSLPYLTKVRFETTAA MARIIGGLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEI			1				IAMCHLRALPGDPSFDAQLGM
AREILVRGMYVRGRKMVAFL CCVALAISAMYELIEWWAALA MGQGADPIGTQGDQWDT0g GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVIDKAWDDLMALONGGVD VMFSNEFSLPYLTKVRFETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEIFT	i			i			NWVIDKAWDDLMALQNGGVD
CCVALAISAMYELIEWWAALA MGQGADDFLGTQGDQWDTQS GYVLRAAWRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSPDAQLGMN WVIDKAWDDLMALQNGGVDV VMFSNEFSLPYLTKVRPETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETTRIAF	l		1				AVMFSNEFSLPYLTKGLVPALV
MGQGADDFLGTQGDQWDTQS GYVLRAA WRINDGDIPRSLSLE PATAL WLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSPDAQLGMN WVIDKA WDDLMALQNGGVD, VMFSNEFSLPYLTKVRFETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHEI	1		1				AREILVRGMYVRGRKMVAFLV
GYÝLRAA WRINDGDIPRSLSLF PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WYIDKA WDDLMALONGGVD. VMFSNESLPLYLTKVRFETTAA MARIIGOLMSDIRIPFGVNVLW DDVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHÉ	1		1				CCVALAISAMYELIEWWAALA
PATALWLDNRI 15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WVUDKAWDDLMALQNGGVDV VMFSNEFSLPYLTKVRPETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETTRHE	1		1				MGQGADDFLGTQGDQWDTQS/
15829 46197 A 15920 I 678 MCHLRALPGDPSFDAQLGMN WYIDKAWDLMALONGGVD/ VMFSNESLPLYLEVRPETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHE			ı		1		GYVLRAAWRINDGDIPRSLSLP
WVIDKAWDDLMALQNGGVD/ VMFSNEFSLPYLTKVREETTAA MARIIGQLMSDIRIPFGVNVLW DPVASFDLAMATGAKFIREIET GAYASDFGVWDTNVGETIRHQ	1		1				
VMFSNEFSLPYLTKV\\(\text{PETTAA}\) MARIIGQLMSDIRIPFGVNVLW\) DPVASFDLAMATGAKFIREIFI\(\text{GAYASDFGVWDTNVGETIRHG}\) GAYASDFGVWDTNVGETIRHG	15829	46197	Λ	15920	I	678	MCHLRALPGDPSFDAQLGMN
MARIGGLMSDIRIPFGYNVLW DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRH			l				WVIDKAWDDLMALQNGGVDA
DPVASFDLAMATGAKFIREIFT GAYASDFGVWDTNVGETIRHO			l				VMFSNEFSLPYLTKVRPETTAA
GAYASDFGVWDTNVGETIRHQ	i		l				MARIIGQLMSDIRIPFGVNVLW
			l				DPVASFDLAMATGAKFIREIFT
HRIGAGEVKTLFNIVPEAAVYL	1		l		l		GAYASDFGVWDTNVGETIRHQ
	1		l				HRIGAGEVKTLFNIVPEAAVYL
GNRDICSIAKSTVFNNHPDALC	1		1		l		GNRDICSIAKSTVFNNHPDALC
VSGLTAGTRTDSALLKRVKET	1		1		l		VSGLTAGTRTDSALLKRVKETV
			1				PDT\VVLANTGVCLENVEEQLSI
ADGCVTATTF			_				
	15830	46198	Α	15921	624	734	TRTLRQIVDSRAIPFT*ERLHSPI
NOPKNDVDESV	1	I	I		l	1	NQPKNPYPPESV

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
					<u> </u>	
15831	46199	Α	15922	I	1923	MPFALGGGDLEETPRSDPGVTF
		ļ				ACSLAQPMRKPTGVLGNRFHD
1						TQEPPVPGGPGLRGFHLWHAPP
		1				LDRPAQFAAIRTDSELMSSANP
						KSFTASPLMTFTPARSRINYRY
1			İ			YPAGTRAKGENEAAFAARIHSF
1	i	l	ĺ			LPYRRPVFWATTMGQAKSAIK
1						PMSTAVTRDASVFQTEYERSRF
		l				GARPWPLRKAPEAMGRGFHEI
			1	ì		GVQCEAPTANTLEPSRRGAVM
		l		l	l	AAGLSFQRTPWREVLGRGEGSF
						LGAGSDPVLRALSLASTPFPGE
		1		ŀ	ŀ	KHVGSWGMGRIVLRSFALASSS
		1		l		LRHSGERIPGTVLPCDDIFACHQ
						RNYAEVFGIWRTTSSVLOPIEPV
		ľ		i		EPKCISQFLYIVLERRSSSSLISLS
				l		FNSASLCRSRHDARYVLQHALF
				i		CFTTNDFGQIFTTFFVSGGSGTR
1					1	MSLPAEFGVIAQSDQKVLFIDA
				i		DLAVVIRITCLPSTSMSWLKEVI
						GTEKAVI/AMCHLRALPGDPSF
						DAQ/LGMNWVIDKAWDDLMA
1		i				LKMVRPETTAAMARI/IGQLMS
1						DIRIPFGVNV/LWDPVASFDLA
1		1				MATGAK/FIREIFTGAYASDFGV
		i		1		/WDTNVARSRRIRASTVFRLAQ
		1				RCOPDASEYGFRMIARVFVDVR
		l				GDONSEASLRVGSGIGPRTCAP
						VAWQSRRFPRLLDRSDDVKRF
						QADTDSLVLH
15832	46200	A	15923	480	900	KTRYSCRGETWSNPQSSQLSAL
1.5552	1.0200	'		1.00	1	IAAVRHPATSNAVCRSARSCCIL
1		1		l		RALRAIOFHTGCSAGNVAPEW
	l			1		CQFAPCWRKPLK\EESDSVFMR
		1		1		VIKESLWOELADITDKTOLEWR
	1			l		EVFQDLNHHGVYHSGEVVGLG
				ŀ		NLVCEKCHFH
15833	46201	A	15924		431	SKVFAVAGCVVLMTTMPDSCN
13833	40201	Ι^	13924	l'	451	SWOKCRNKTMNKKIHSLALLV
				ł		,
1		1		I		NLGIYGVAQAQEPTDTQAPGVS
1		1		1		TITADEIRKNPVARDVSKIIRTM
1		ı	İ			PGVNLTGNSTSGQRGNNRQIDI
1						RGMGPENTLIL/IDGKPVSSRNS
		-			10.00	VRQGYRLECGDG
15834	46202	В	15925	1	1352	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15835	46203	Α	15926	31	546	CYCILFVSFPSNRPLCCRSAGVC
						ORYTPDPFCLGITSRGCRRAKIT
		ŀ				ACSFLWKLHPRGALARCOPELS
		ŀ				CWRCLST\LLEGVSPSGGTGVR
				ļ.		DPYEEAVCPLAELKCCAGRSA
						VLFRAGROGRLSLLKLFPOLPF
		l				SPCVLCLSKLVPVSPNSTRTSKL
				•		LSIPYSRGPALLEVLS
15836	46204	A	15927	2	479	
15837	46205	Α	15928	1082	1386	ADPFQEREAEKYANPIPSREFIL
1				1		EHLTKREKPASRDELAVELHIE
			ŀ	l		GEEQLEGLRR/RQCYALPERLD
İ						LVKGTVIGHRDGYGFLRVEGR
				1		KDDLYLS\SEQMKTC
15838	46206	Α	15929	3	40I	
15839	46207	В	15930	1	3675	
15840	46208	Α	15931	1749	1941	SQIPTSEAQVMSNLMGTNGVLR
1					•	PARAYRNEAVLS*ALVSAMYS
						QTFRNASFGVPQMFSTTSGV
I584I	46209	Α	15932	5	1465	KRSTTAQIRWISPLTSLKPRSKR
						QSKSCATKPKKWYVAAPCCWC
						SPTGISLKTRLPVPAPMAVGAIQ
				l		TRLVDQSLRCDANIIVETASAR
				l		DPHHFAVLLGFGATAIYPYLAY
						ETLGRLVDTHAIAKDYRTVML
						NYRNGINKGLYKIMSKMGISTI
						ASYRCSKLFEAVGLHDDVVGL
ĺ						CFQGAVSRIGGASFEDFQQDLL
	t					NLSKRAWWRCRLSATIGCCDC
						TRGYLGGARITGQISSITVWGVI
						IPVVGLCIIGWFWFSPTLYVDS
						WNPHHAPFFSAVGSSIAMTLW
						AFLGLESACANTDVVENPERN
						VPIAVLGGTLGAAVIYIVSTNVI
						AGIVPNMELANSTAPFGLAFAQ
l		1				MFTPEVGKVIMA\LMEMSCAGS
		1				ILGGKFTIAQVFKSSSDEGYFLT
						IFSRVTKVDAPVQGMLTIVIIQS
		1	1			ELALMTISPSLNSQLKEMVNLA
						VVTNIIPYILSMAALVIIQKVAN
		乚				VPPSKAKVANFVAFVGARNSF
15842	46210	A	15933	I	3666	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15843	46211	Α	15934	979	1626	FSATITPPFIALQRVRFHPLDAL
						LLFVISLWLQLGLPDELSRFTPF
						CLALPIIALAWHYGWQGALIAT
		1				LMNAIALIASQTGLLLGAGIQR
						LRELNQSLQKELARNQHLAERL
						LETEESVRRDVARELHDDIGQT
		1				ITAIRTQAGIVQRPGAGHPLTDA
						GNGAGRARYCQPSRMANR*IS
		l				VKRKVVEHRCFVDFTFTNDDG
						AVNVDRRAVRGVFIAAS
15844	46212	С	15935	1	666	\
15845	46213	Α	15936	I	1137	MGKFGHTEIGEVDVRDWSDAL
1						QAKELQKLPGHSRKLEDPPTSL
						FMLLNPHSDQFSSNAHHHLHA
						TAIDDDYTDPTKSTYALGLNWS
						NSILGGFNISGYKTYSYDGDND
						SSNLNINWNKAFKHATVSVNW
						QHQLSASENNEDDGDLFYVNIS
						IPFGRSNTATLYTRHDDHKTHY
		l		ļ		GTGVMGVVSDEMSYYVNAER
	İ					DHDEPTLPAQNKAQEVLLDVV
						LDEAKIGVASMLGSRVRVKTW
1				ŀ		SWFADDKQEIRQGGFAGWLTD
		l				GTPLWVTGSGTSKTVLTRYAT
ļ		1				VLNRVLPVPTQVASGQCVEVE
1		l		ŀ		LFARYPLKKITAEKSTTAVNPG
				ŀ		VLNGRYRVTFTNGNHITFVSHG
		l				ETTLLSEKGKLKLQSHLDREEY
	L	_				VARVLDREAK\STPP
15846	46214	Α	15937	323	695	FTGPRRSLEIEVAELKQRLDSLL
						AIGAGGSVCWTLQGAWSPTRA
						KAGELKTQLATAASLR/MDKH
	1		i	1		RSNSVGPHDLYFTLLDDYLHV
1						VDTALWLSGGKASLDGGTLLT NDAGEMLFAEHHFSAGPL
15847	46215	١.	15938	I	235	NDAGEMLFAEHHFSAGPL
15847	46215	A	15938	98	593	MSKVRLMRNLSPAPPNAVAKG
13848	46210	A	13939	98	393	ESKPGIMSGWVF*AEWAYLITG
						*AKKKVYGSS/ARIHLRALRYY
1		1				SNWROYFAGYTFGROYWOSPE
		l		ļ		DDHLPLLREFLARHEYMQKYL
		1				LGTESGIQGEELGASDGIKPEEV
		l	l	1		EWQTAAIEGKLDLLVTLDFRM
		1	1			SSTCLFSDIVLPTAT
15849	46217	A	15940	367	677	LSSCLRATQSAPAVRNWKARFL
13849	1021/	l^	13940	J**	l°′′	RCTVVVWQKRSSTGRNMYLPV
		1				RPTILKLRPTWHVTW*LSGASL
1						RNWVHCCTRKKKVKCSSAVA*
1		1				RKRNICPMKLHVSSTRK
			<u> </u>	L	l	KKKINICTIVIKLEI VOOTKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15850	46218	Ā	15941	1	921	MEMSSLEFLHSICSHLKCHLLY
13030	10210	ľ.,	1.57	ľ	12.	YIASAYNYDWYTOPAKLSFISE
i		l				GEIKYFTDKOMLRDFVTTRPAL
İ			İ			KELLKEAVNMERRNRWVDIGP
						GTELDCISQAYAQFVASTAECC
						GEGGIKAWDYVRMGFLSRMG
						VLNNWLTRRRKLLWSARSRIH
						LRALRYYSNWROYFAGYTFGR
		l	İ			QYWQSPEDDHLPLLREFLARYE
						RPMKKLAGRSSIAILAFRGHDQ
		l				RDYEFPSKSALNIN*LRWALAS
						VK*TTACATGVFPVSVTGARRF
15851	46219	Α	15942	531	637	SIFARLNASRRC*TSALYALLVR
						LPIIQITAKGIL
15852	46220	В	15943	120	881	
15853	46221	Α	15944	1	1583	MADRVDSKCSRSRPGAVAHNY
		1				NPSTLGGRGGRITFWLLSANPSI
		1				NRGTLEKESRTVAQRLSVLHGI
						NAPEFFDKAVFSSLVLTLRDEG
						YISDSGDAEPAETMKVYQLLAE
		l				LITSDLMIFNISATVLYITAFIAC
						SAAVDLTSLRGTRPYNQRAAA
						SKEIEDLKKKLTEGEEISGCDVS
						GSDGDDNEEGKIGEDEEKRNT
						GGNKVFPERTVETPATADAER
						KALETKLDVEEETRLQLNSRNR
						KKDLKTRQEHQSLPEKFSGLQK
						NVLLEAVIDPALIAQLRQAYEE
						AL\RIAQAPAKSVWQPEEYEEA
						EHEILLAFRALLASDSERFLPSA
1						WQRFIQQLNYCSMEEIDELRWS
						LCTIAMNTAHLSFECVVLLAER
						LRWLQEENTGEIDEEELESFLY
						AIAKGNVFNFQTILHLPVAVQN
l		1	1	l		DTIDFYQMFARIWLPGAPGAYC
				i		DGLHTSKRTTCTTSTHQRQRKD
1		l		l		TPPALNPHTATCTPPHTWLGAP
				1		PPASMRSGPHNAPTPPFTGSTPR
L				L		HTHDHTNITNANIPHNVNSN

SEQ ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Uuknown,
NO:	of peptide	hod	in USSN	Incation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15854	46222	Α	15945	1	1228	MGVLNNWLSEEESLWIQSRIHL
		l		l		RALRYYSNWRQYFAGYTFGRQ
						YWQSPEDDHLPLLREFLARYER
ľ						PMKNCWKILDIEETTDVDIIRRA
				l		YLALLPSFHPETDPQDFKQLRQ
						AYEEALRI\AQSPAKSVWQPEE
						YEVAEHEIL\LAYRALLASDSER
						FLPSAWQRFIQQLNYCSMEEID
1		1			1	ELRWSLCTIAMNTAHLSFECVV
1		l				LLAERLRWLQEENTGEIDEEEL
						GSFLYAIAKGNVFNFQTILHLPV
	1					ARRPDICTOHNDOSRCEGYOIL
1		l				SDKRGHOHGGGITALYOCGNA
1		l				DPRAKGERLFFDAPAKDSTQTR
						AEDAHYAGSNNMSPPDQQGDG
		1				RKQIWESVRVHTPVRCQQRSLP
			1			VTSYLNGGEIFNLIGLICRTRRIA
		l				ARWVKSHRNRARSFECLKRHR
ı		1				FVVAFNRRKPGLV
15855	46223	A	15946	71	323	AAGAVVSAMPKAKGKTRROK
13833	40223	<u>۱</u> ^	13940	l''	323	FGYSVNRKRLNRNARRKAA\R
1		1			İ	GSNDLEAEASLPEKKGNTLSRD
1						LIDYVRYMVENHGEDYKVSGS
15856	46224	A	15947	48	627	AAGAVVSAMPKAKGKTRROK\
15856	46224	I ^A	13947	48	027	FGYSVNRKR\MSRNAKRKAAR\
1	1		1			
						GLECSHIRHAWDHAKSVRQNL
						AEMGLA\VDPNRA\VPSRKRKV
						KAME\VDIEEEA*KNLYRKPYV
						LNDLEAEASLSEKKGNTL\SRD
	1					LHLTMYRYMV\ENHGE\DYKAI
		l				A\RD\EKNYY\QDTPKQI\RS\KIN
i				i		VYKRFYPAEWQDFLDSLQKRK
	-					MEVE
15857	46225	Α	15948	796	1301	LTKKDYISPNARKAFVQRLPER
1				I		NLLTRGLKRQRQPCLSCWCGG
1		1				QARMPKAKG\KTRRQKFGYSV
						NRKRLNRNARRKAAPRIE/CVR
		1				GLGFRVAVGLGRDGPRDDGSA
1		1				MSLFAGSSHIRHAWDHAKSVR
1		1		1		QNLAEMGLAVDPNMAVPLRK
				I		RKVLTWQGLGPCLLYSSPAILT
15858	46226	Α	15949	512	675	SQDFGPPSAAPSPSGMP\PGPSG
		1				CSSHRQQEDPPDGAAAPPGHGL
						RPGRLQELLV
15859	46227	В	15950	1	717	· ·
11.000	1	_		L	L	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15860	46228	A	15951	587	1387	RARSRILGNPLRPLAPAPSNDV
						GVQRA\FRPIRCRKAQPVPGSV
						ANRRILROHTRPDPLGTNRSEP
						GWWDRSWSPDQAKSGQSCVA
						KIKNKNHMIISTDSEKAFDKIOH
						RFMIKTLSKISIQGTHF\KIIKAIC
						DKLTANIILNGEKLKAFPLRTE
						MROGCPRLSLLFNIVLEVLAKA
						IRQEKEIKDIQIAKEEVRLSLFA
						DDMIVYLENPKDSFRKLLELIK
						EFSKVSGYKIKNOYCENDHTA
						VSGVGSFQWICGLADVKNEAA
15861	46229	A	15952	3	3201 ·	SERGFRAGDLRYPTHFLSTNSV
13801	40229	l ^A	13932	3	3201	LASVTASLKEHPRGTLLSDGSP
1						ALSRNVGMTVSOKGGPOPTPSP
1				ļ		AGPGTOLGPITGEMDEADSAFL
				ł		KFKQTADDSLSLTSPNTESIFVE
						DPYTASLRSEIESDGHEFEAES
						WSLAVDAAYAKKQKREVVKR
				ļ		QDVLYELMQTEVHHVRTLKIM
						LKVYSRALQEVL\QFISTAIGRL
1						FPCADDLLETHSHFLARLKERR
						QESLEEGSDRNYVIQKIGDLLV
		_				QQFSGENGDRLIEK
15862	46230	В	15953	83	1969	
15863	46231	Α	15954	1	444	SSESAVEFSEPASGSSPSWELGS
						EGAIRDKKTSLGAHVGLSALGR
						ALENTIAFQGKETDVQKSYDSP
						QGDCL\QDVKGKRPTLE*PSLE
						NPETRRFLDASYAWRESCLSW
						HHHTAHDIFAGKPVDPLPGSTR
		_				GQDSAVCSQGQFLIVR
15864	46232	Α	15955	2	236	SEGAKRDKKTSLGAHVVLSAL
1		l	l	l		GRALEKTIAFQGSS/APSKGRKL
1						RSRKVMTRPKGIAWQDVKGKR
						PTLE*PSLENPEVR
15865	46233	Α	15956	1	459	KKVGNYYTTPIYRFRMKCHLC
1		l				VNYIEMQTDPANCDYVIVSGA
		l		l		QRKEERWDMADNEQVLTTGE
		l	1	l		RHPLTCLGAL/DPESALGPPKPS
1		l		l		RALIVAEHEKKQKLETDAMFR
				l		LEHGEADRSTLKKALAHT\DHI
				1		QEAQSAWKDDFALNSMLRRRF
		l		l		REKKKA

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15866	46234	A	15957	564	1189	AAPKMGERKGVNKYYPPDFNP
15866	46234	Α.	13937	364	1189	EEAWLSSTDYHNSHPFRERASG
		l				SCSQGI\LIIRFEMPYNIWCEG\C
		j				
		1				RTNFGMGVRYNAEKKKVGNY
l				1		YTTPIYRFRMKCHLCVNYIEMQ
						TDPAN\FDYVIVSGAQRKEERL
1		1				GHWRTMNRLLTHRGIEEEAEA
						WKTDAMFRAGTMGRPTAAHF
		l				KKALPKLEPHSGRPRAPWKDD
						FALNSMLRRRFRVPSKP
15867	46235	Α	15958	2	1047	FVDGKLHGRGSTDDIGPVAGWI
						NALEAYQKTGQETPVNVRFCL
		1		l		EGMEESGSEGLDELIFARKDTF
						FKDVDCVCISDNYWLGKKKPCI
ļ						TYGLRGICYFFIEVECSNKDLHS
						GVYGGSVHEAMTDLILLMGSL
ŀ		ł				VDKRGNILIPGINEAVAAVTEEE
1				İ		HKLYDDIDFDIKEFAKDVGAQI
						LLHSHKKDILMHRWRYPSLSLH
		l				GIEGAFSGSGAKTVIPRKVVGK
1		ı				FSIRLVPNMTPEVVGEQVTSYL
		l				TKKFAELRSPNEFKVYMGHGG
i						KPWVSDFSHP\HYLAGRRAMK
						TVFGVEPDLTREGGSIPVTLTFQ
		1		i		EATGKNVMLLPVGSADDGAHS
1		l		1		QNEKLNRYNYIEGTKMLAAYL
15868	46236	Α	15959	163	1388	RLVERWRPSGTLFKYIDENQDR
		l				YIKKLAKWVAIQSVSAWPEKR
		l				GEIRRMMEVAAADVKQLGGSV
		l				ELVDIGKOKLPDGSEIPLPPILL\
		İ				GRLGSDPOMKTVCIYGHLDVO
		l				PAALEDGWDSEPFTLVERDGLT
		1		1		WCSSCRGPYCRGSS/DRLSWKL
		1				PSVMWGDSREPHLARLD\HPSG
		l				LEPORPGSTWGGGDDRRGGRA
		l		1		LSROLPYAGYAAAOCKGEVRP
i		l				OVSSSRLVVECSNKDLHSGVY
		l				GGSVHEAMTDLILLMEEHKLY
		1		I		DDIDFDIEEFAKDVGAQILLHSH
		1		1	1	KSHLHLDLLPVVVRLLGQALFH
		l		1		TAHFPDNIPSSSKDILMHRWRY
				i		PSLSLHGIEGAFSGSGAKTVIPR
		1				KVVGKFSIRLVPNMTPEVVGEO
1		1				ACGAGTRESMSSLGYPSRAED
		1			1	
	<u> </u>				L	DSGLSALPSQPQPFILYAT

SEQ ID NO:	SEQ ID NO: nf peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15869	46237	А	15960	3	433	FETLFSQVQLFISTCNGEHIRYA TDTFAGLCIQLTNALVERKQPL RGIGILKQAIDKMQMNTNQLTS IHADLCQAITTPAMAVSHIMLE SYKKYILVSLILLGKVQQLPKY TSP/QLVGRFIKPLSNA\YHELR HKLIVSTNIPC
15870	46238	A	15961	46	1481	TSGSDVAAREDLRWIAAAPPPR AAAFSRACCLRRSGRPGENMA SALEQFVMSVRLLLSQGGSCFP GQMTQLCELIMMSGELLAKNLS ALEQFVMSVRLLLSQGGSCFP GQMTQLCELIMMSGELLAKNLS LLDTVLGALDDVQEHSLGVLA VLEVKFSMPSVPDFETLFSQVQ LEISTCNGEHIRYATDTWQETV LLKSMVLVLVYJYQPLRGIGIL KQAIDKMQMNTNQLTSHADL CQLCLLAKCKFAALPYLDGDM MDICKENGAYDAKHFLCYYYYQ AITTPALMAVRHILWWESYKKY ILVSLILLGKVQQLPKIFHLKLY GMIYTGLKNFERALYFYGV GYPLSLLSNAYHELAQVYSTN NPSELRNLVNKHSETFTRDNN NGLVKQCLSSLYKKNIQRLTKT FLTLSLQDMASRVQLSGPQEAE KYYLHMIEDGEIFSNIQKDGM VSYFHDNPEKYNNPAMLHNIDQ EMLKCIELDERLKAMQGITVN
15871	46239	В	15962	1	3411	
15872	46240	Α	15963	689	925	
15873	46241	Α	15964	I	244	GSAPQSCASPAPVSRSAAPGTA EAAEPWAIVPGVEQKNYYVCV PLGLIGNCTVTSSG\\SLDCADW CLKWKSILMESLEVP
15874	46242	Α	15965	469	604	
15875	46243	A	15966	286	433	
15876	46244	A	15967	4356	4643	DCPLARNKSKPQTVCLKARQTF QAINFSRALRITGYLTSFPSLSRS TLPSSAGDFRASAWA*SGPTV* TLTSVPSVGINSSYIYEPVFRTLL SGE
15877	46245	В	15968	1	1472	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15878	46246	A	15969	918	1619	KITHULSWGSPKSTWKAGWLQ KOKTTOEKKAESKAPIEEPAG AGLNKDKETEERSDGERVAEV APEERENVAVAKLQESQFONA VSCSGGRRKEDEREDWGKRTQ RTHPPAAAQAPGQCPPGSTEVG AGGGLRWTPSPGRQPAGVAAD MGITVLTVPLQRCLQNSWAKE PVLVVSFVIGSLTVILPPLSPYTK YSIMINEATPHYPYPVPHDDON
		l				M/PDMCSHPQGPQGPSLEWL
15879	46247	В	15970	183	459	
15880	46248	A	15971	864	1197	GSFRLGHGRIPPFHLRGMKESL GPDFGDPCPVCPLNSGPGPP\GV SAGGAGAGCGGMGRRDAWVP TGPVAESPGGHLAPISPGRQGA PWGGPCGQLGAGTLGQAVCG MTQDP
15881	46249	Α	15972	1	441	
15882	46250	Α	15973	3	182	
15883	46251	Α	15974	23	475	PLEKGRQFYPQYEDQITIFTDYL EEYPDTDELVWILWKQHLLKT EKSKLLSDISARLWFTYRRKFSP /ILGGTGPSSDAG\WGCMLRCG QMMLA\QAL\ICRHLGRDWSW EKQIEQPQEYQRILQCFLNRKD CCYSIHQMAQMGVGEGKSIGE
15884	46252	Α	15975	224	418	
15885	46253	A	15976	743	1220	WCWGHRVRLVLAAGPSSRGVL LLLVVLPSWASFLVRVCAWVG VLGDDGVLGGFLLWLGVVGRP LAVLRADLAVCVGVVCACVPF VVLPICAALVRVGCSLVEAALG \LGARPLKTFFAVVVPLTKGGV VAGSMLVFVPAVGEFVIPELLG GPDSIVIGRV
15886	46254	A	15977	3	3492	RKHKVGILYCKPÖQSSEEMSN NEEAGPAFEKFLSLIGEKVCLK GTTKYAAQLDVKTDSTGTHSL YTMYQDYEIMFHVSTLLPYTFN NROQULRKRHIGHDIVTILPOEP GALPFTPKNIRSHFQHVFIIVRV HNPCTDNVCYSMAVTRSKDAP PFGPIPPSGTTFKSDVFRADLA AKVI*RLRNAAEQVPTSFTPWA TRTRQEYLKDLAENCVSNTPID STGKFNLISLTSKKKEKTKARA GAEQHSAGAIA
15887	46255	В	15978	402	2884	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	ol peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, *possible nucleotide insertion)
15888	46256	Α	15979	294	1721	AVRPGRHDPSAGLRSAAAASQ*
				l		VRGHPGGLCA*GSIQGPGASAA
	i i	1		l		EGPPAHSRAA*SKGGAELRTR*
		l		1		AGGRRPAGPHAQRP*LP*GPRV
		l		l		PLRPPSGTFLLP/PPACTPITSRPS
		l		l		*KSCSRASTPPTWSS*IPASGTSF
	l		1	1	i	GMVRTPGEA\PGEPGEPVPVPG
		1				PGAARVLPPGVE/PRPCLWARK
	1			1		SPGVFFRPSSGGRRPPS*KTKWS
	1	l		Ì		KPTSTAPPR\P*T*LRCTGLAFPL
		1		l		PPREGEPALGRGALEWTCAPLP
	1				LPAAAGPRGRRLGCGAAPPPPR	
					GRVDQEEK\PGPRVEGPPQANR	
	l				NHPALPLSPPLPSPT/*PPPAWVP	
						TPALAAAPAPVPTA\$SSHSPSPG
	İ	1				NAPVPTGSPRCLFFLRPYFPVGS
					ILLPFRCPLISPCRFLRRRQFYG	
					WSSAAYALLPHTPLSAACPSGP	
		1		1		TTALHGQPQSSCSCCQEATFTV
	1	l				KSGVVQSGRPSLRQFYVHDFCK
	1	l				IQPQQLLVSHATTKATTDRYST
		1				ASCVAR
15889	46257	В	15980	35	314	
15890	46258	A	15981	1	197	
15891	46259	A	15982	3	3889	
15892	46260	В	15983	172	287	
15893	46261	Α	15984	8	166	
15894	46262	Α	15985	250	1095	
15895	46263	A	15986	366	870	NHSLAKKGTAHGLGMEKCCGR
	i			I		SPFFSFFPSWIAKPTLRSSSSWPH
				l		SSSSPSMSSPLQRAVGDT\KRAL
						SASSSSSASLPFDDRDSNHPSEA
						CTTPGGGWEVADGNTDSLLAD
						EGSDFEDSFNRNVKKKAAKRPF
						KTTPV/AKQPKKGSRVVHRHSR
		L_				KQSEPPANDLFNAV
15896	46264	Α	15987	30	229	ALETNIGSSYNYGSEDQAEFLC
		1	1	l		VVSKELHSTPNGLSSESSRKTK\
						CYKPEAPGCEAPDHLQGLGVPI
15897	46265	С	15988	1	930	
15898	46266	Α	15989	1	609	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,217	sequence	or peptide sequence	decidon, (-possible nucleotide insertion)
15899	46267	Α	15990	1	875	MRMEAGEAAPPAGAGGRAAG
						GWGKWVRLNVGGTVFLTTRQ
						TLCREQKSFLSRLCQGEELOSD
				Ì		RDETGAYLIDRDPTYFGPILNFL
		ŀ		}		RHGKLVLDKDMAEEGVLEEAE
		l		İ		FYNIGPLIRIIKDRMEEKDYTVT
1						Q\VPPKHVYRVL\QCQEEELTQ
1				l .		MVSTMSDGWRFEQLVNIGSSY
İ				ł		NYGSEDQAEFLCVVSKELHSTP
				[i	NGLSSESSRKTKSTEEQLEEQQ
		l		Ì		QQEEEVEEVEVEQVQVEADAQ
		l l				EKGSRPHPLRPEAELAVRASPR
		1		Į.		PLARPQSCHPCCYKPEEPGCEA
		1				PDHLQGLGVPI
15900	46268	Α	15991	1	2448	MRRGETRSRRPPQCWAFAHVV
		l				SRSSRRGEFFSFDRWKKHRSEV
1				l		RESAQVPTTKREPVLFQHKINIK
1		l				IVIALVDILFALGLGNLVCVSPV
1		l		1		QLFSVQTDSRATGDQRLPSWT
		1				AQRYGTRTLGPDGFTERCPYGP
	1	l		Ì		SPESLLCSALPWLRDLGMPVAE
İ		l				APQVAGGQGDGGDGEEAEPEG
İ		l				MFKACEDSKRKARGYLRLVPL
		ļ				FVLLALLVLASAGLEMLSDPGS
1		1		İ	i	RSYGSGLLPAWAKCPKAPHQR
		1		l		SLIPPDYAPGVNHPPPQERCSM
1		1		ŀ		GLRTQGTLGRHRVLSAQQVAF
				1		SVSLLMAKIAFAKQKSDDIIPDF
		1		l		KLFMVTLVHRPNRLSARVLTM
		l		ł		VPVLPAPCPMLLPLLEGYKAEV
		1		ł		MVSQVYSGSLRVLNRHFSQDL
1		1		i		TRRESSAFRSETAKAQKMLKEL
1		l	ł	l		ITSTRLGTYYNSSSVYSFGEGPL
		l	l	l		TCFFWFILQIPEHRRLMLSPEVV
		l		l .		QALLVEELLSTVNSSAAVPYRA
		l		}		EYEVDPEGLVILGQYCEWKPPG
1		l				RRMQTPRPAMRMEAGEAAPPA
			l			GAGGRAAGGWGKWVRLNVG
1		l		1		GTVFLTTRQTLCREQKSFLSRL
		l	l			CQGEELQSDRDETGAYLIDRDP TYFGPILNFLRHGKLVLDKDMA
1		l				
		1	1	l		EEGSQVWTVTSGGWGDRNAV
						LTGMGRALSLFLWLIQYLKFCL
		1	l	1		VRAWVLEEAEFYNIGPLIRIIKD
						RMEEKDYTVTQVPPKHVYRVL
		L_			l	QCQEEELTQMVSTMSDGWRFE

SEO ID	SEO ID NO:	Mct	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminu acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15901	46269	Α	15992	li -	573	MPGLLSRPRRGSLRARPCGGGE
		ŀ				ALPEHVAIDVCPGPIRPIQQISG
						YFPHFPRDLPHDAPARPATASA
			1		i	GRRRPSDGARDHDKDGDHLFG
		1				SLTIQHRPW/PAPSPDFPFTPFAV
			ŀ			LYPG\LKPMDHNGLADPCVKL
		l				HLRPGARKVPEACQLAPDLQD
			l			SSCLTAITKAHDVLTVFLPAAL
l	i					KIQGNKYLTFIECQKLF
15902	46270	Α	15993	500	2022	WTREKEPPGVETHGKASRTTL
		l			i	GLSLWCCLHLSEIQSSFISDCAR
			l			PRCSGRLPGSGPMLLATPHVLA
l						TVVASSCSKQLWWLPPKVATD
						QQQMKLKGLFIAGAPPGSPR/G
						VQPHLQPIRNMSVSRTMEDSCE
						LDLVYVTERIIAVSFPSTANEEN
İ						FRSNLREVAQMLKSKHGGNYL
l						LFNLSERRPDITKLHAKVLEFG
						WPDLHTPALEKICSICKAMDTW
		l				LNADPHNVVVLHNKGNRGRIG
		l				VVIAAYMHYSNISASADQALD
						RFAMKRFYEDKIVPIGQPSQRR
		ł	1			YVHYFSGLLSGSIKMNNKPLFL
						HHVIMHGIPNFESKGGCRPFLRI
						YQAMQPVYTSGIYNIPGDSQTS
ĺ		l				VCITIEPGLLLKGDILNSEILFGV
						DELSLTGFGLKCYHKKFRSPAR
		l				DVIFRVQFHTCAIHDLGVVFGK
						EDLDDAFKDDRFPEYGKVEFVF
						SYGPEKIQGMEHLENGPSVSVD
				1		YNTSDPLIRWDSYDNFSGHRDD
						GMEGAQHPQIGQDTPPGPSDPP
15903	46271	Α	15994	171	720	LSGSSÄGKVAAPCVPPSNHELV
					1	PITTENAPKNVLDKGERPSRGG
		l				NTRKSLE\TTASTGSPR/GVQPH
ŀ		l				LQPIINMNVSRTMEDSCELGPV
		l				YRTERNIAVSFPSTANEENFRSN
l		1	1			LREVAQMLKSKHGGNYLQRN
i		1	l	l	1	VSL/RRLSPSLRHPGPWPPSQEP
				l		GMCQGEPGPPATLWPMPCGPT
		_				VLESPGGPGP
15904	46272	В	15995	145	972	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequenee		
15905	46273	A	15996	2	1096	WPVLTAAGCWLQDPAGHRGG
13903	40273	^	13990	1	1090	RAGGAGRAVLFQLQAGRAELR
1	1	1	1			
						GLRARGWARSWAPAPGCQRTG
						RDAAAAGRRAGRAGALLGFTC
				1		LLLAALCGHLGPSWRGGLAPG
						GEWIWLRASETRGPSGPRPALM
ļ.						GSQKIRNITRTRHGPLHWLHSG
					·	CTTATPFCLTRSCSDAACSLKG
						LPCLCPCGLSVLIRVRLELRRHQ
ŀ						RAGHTIPRIFQAVVQRQPERLA
			ļ			LVDAGTGECWTFAQLDAYSNA
1			l			VANLFRQLGFAPGDVVAIFL/E
						GRPEFVGLWLGLAKAG/MEAA
						LLNVNLRREPLAF/CLGTSGAK
l						ALIFGGEMVRAPPPNAQAPPLT
						PVSPADRLFYIYTSGTTGLPKA
		1				AIVVHSSARPSCPSSRTLRAAHR
15906	46274	Α	15997	397	597	SIKSPEIHSPSTRIRMGCSSLPTE
1						EAHAEGVIELSRIKCVEIVKSD/
1	ł					DRIPCHYKYPFQVSPSGGSSP
15907	46275	Α	15998	1922	3886	EGDNKLRPRIRRKLFPLVVLRG
1			l			DAQGAPPFKNWIMNNFILLEEQ
		l				LIKKSQQKRRTSPSNFKVRFFVL
						TKASLAYFEDRHGKKRT\RKGS
		1				IELSRIKCVEIVKSDISIPCHYKY
1		1				PFQVVHDNYLLYVFAPDRESR
1	1					QRWVLALKEETRNNNSLVPKY
1	İ	1		i i		HPNFWMDGKWRCCSQLEKLA
ı						TGCAQYDPTKNASKKPLPPTPE
	1					DNRRPLWEPEETVVIALYDYQT
		l				NDPOELALRRNEEYCLLDSSEI
	1					HWWRVQDRNGHEGYVPSSYL
	İ					VEKSPNNLETYEWYNKSISRDK
						AEKLLLDTGKEGAFMVRDSRT
1						AGTYTVSVFTKAVVSENNPCIK
1						HYHIKETNDNPKRYYVAEKYV
1						FDSIPLLINYHQHNGGGLVTRL
1						RYPVCFGRQKAPVTAGLRYGK
						WVIDPSELTFVQEIGSGQFGLV
		ı				HLGYWLNKDKVAIKTIREGAM
ľ	1	ł				SEEDFIEEAEVMMKLSHPKLVQ
Į.		ļ				LYGVCLEQAPICLVFEFMEHGC
		1				LSDYLRTQRGLFAAETLLGMCL
1		1		ŀ		DVCEGMAYLEEACVIHRDLAA
		1				RNCLVGENQVIKVSDFGMTRF
1		1				VLDDQYTSSTGTKFPVKWASPE
		1				VFSFSRYSSKSDVWSFGVLMW
1		1		1		
		1		1		EVFSEGKIPYENRSNSEVVEDIS
		1				TGFRLYKPRLASTHVYQIMNHC
15005	14000	-	1.5000	-	1506	WKERPEDRPAFSRLLRQLAEIA
15908	46276	В	15999	58	1526	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
15909	46277	Α	16000	3	378	GASPDLGGRVLGAGASAESPR
						GRPFGPSPLGGPPVPGCQCAAP
				l		SLRDPWGLWAPVLQMTGSYEF
	į	l		ŀ		KLNQPPEDGISSVKLSPHTLPSP
						GLSPSWET\SVRLYDV\PGNFM
						AVKYQ\HTGARPGNGPF
15910	46278	В	16001	83	2851	
15911	46279	Α	16002	296	1186	AENHPGP\LVVEGKMTAPERM
						KHESNYLRGTIAE\NLNDGLTG
						GFKGANFLLIRFHGMYQQDDR
						DIRAERAEQKLEPRHAMLLRCR
		ļ				LPGGVITTKQWQAIDKFAGENT
						IYGSIRLTNRQTFQFHGILKKNV
						KPVHQMLHSVGLDALATAND
		1				MNRNVLCTSNPYESQLHAEAY
		l				EWAKKISKQLLPRTRAYAEIWL
		i		1		DQEKVATTDEEPILGQTYLPRK
		l				FKTTVVIPPQNDIDLHANDMNF
		1				VAIAENGKLVGFNLLVGGGLSI
		1	l			EHGNKKTYARTASEFGYLPLEH
						TLAVAEAVVTTQRDWG
15912	46280	Α	16003	225	386	GLCIPVVRSPCQLWGQKGPCM
		1				LRDEGRVFT*LSTKKNCPGNMR
						RKYETVDCRR
15913	46281	A	16004	386	739	PLLIHWRFATIEEADDHRLAAG
						ARTFQ/PLCGPPQAQAVWR*AQ
ļ						*FQFADRFPASRWRRASSGRRF
		l	1			SRSGRARRCQSVAKRRRQIHEC
		ı				GRSTPDCRCRMRQQYRCCRDA
						RRYQNRAPRR
15914	46282	В	16005	10	1468	
15915	46283	Α	16006	3	1267	GRSRR**PPSAVPHD\PAVQRER
		1				DTQSLSVII*MQIYNFSNTTHLT
		l				NHNTFLTFHKTFTAETFQHIGP
	1	1				QPTTDTPYHTAEAYCCQLVKLI
						RRQALRRCWQGAAILKCNSKP
l	l	l			l	TNHCTS

sequence 09/540,217 eulon for peptide of peptide sequence deteiton, \square period of peptide sequence deteitor, riod of period of	n, /=possible nucleotide sssible nucleotide insertion) FRALSRLPDLWHRSI
sequence 09/540,217 culon for peptide of peptide sequence deterion, \(\sigma_p \) 15916 46284 A 16007 I 1673 MTPICTV. GTLTEAL SYSGLVE. SYSGLVE ESGGLVEFI	
15916 46284 A 16007 I 1673 MTPICTVI GTLTEAL SYSGLYG ESGIWFFI	FRALSRLPDLWHRSI
GTLTEAL SYSGLYG ESGIWPFI	FRALSRLPDLWHRSI
GTLTEAL SYSGLYG ESGIWPFI	
SYSGLYG ESGIWPFI	POTDKROEOIAVALT
ESGIWPFI	LGYGSKEIOLTYSGT
	PYSPNATHKGCGRHR
I I I I I I I I I I I I I I I I I I I	IFLVIEPOSALGSCPSP
	CVFGAIRLLVLRDAP
SFSRLLN°	TLASGESLOGGPGFV
	AFLVVRNPTGTPMFT
	VPTFPIPPVFGRLGSF
VLOLOGE	RSPGQCEYDRELVSG
	LLVLGTLHLLSYKDA
GVDIDAC	NALVGRIKGVVKKT
	GGLGGFGALCALPOK
	SGTDGVGTRRRLAL
	IGIDLVAMCGNDLVV
	FLDYYATGKLDVDT
	IAEEKSEILDGSKVSD
	GSSGLHSNGYSLVRQ
	AADDVLSDADAVSW
	SGLRKKINQGAVSFV
	AOMVNYGFFGDIDV
	LAPDGRVWLTSGIRN
	AKKVIIELNHYHDPR
	VIPGAASRRNNGSIFH
AMDRLG!	NPHVQIDPEK\IVAVV
15917 46285 A 16008 425 595	
15918 46286 A 16009 315 721 QCFRALG	GHRRIADRGENIQV
	NDLLKWLNALDEKY
RSRLVW.	ARSSKNSLCASEANW
LEMNNRI	LSKGDLPCIRPILIIMLI
R*TCSYG	FTALR*TQNAAGGGD
SLAGGIG	IFTLVFFTLPAYAIRW
LPR	
15919 46287 B 16010 652 858	
15920 46288 A 16011 238 2526	
	KTSGAGLSITSEQRR
AAVSRIS	VTSSLLASAG/VRYRQ
	/RERPVDKL/PR**TPC
	GPSR*WW/MARVLIF
	RIVTVSPMRIGFSNRII
RPEIKLAN	KISCKPKPRPTPSAAT
SHCNFDH	ISIPIIEKPTSPPISISRY
LVMVVM	Α
15922 46290 A 16013 2 527	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			i	sequence		
15923	46291	A	16014	1	1734	MVNGCDSTNASSYANVNGCDS
13723	40271	^	10014	ľ	1734	TNASSYANGERCDSTNASSYA
		l				NGQRVRSTNASSYANGERVRL
				l		DNASSYAIGERVRLDERTSYAN
		l		l		GERVRLDERILLANGDRVRHHE
				l		RILERKATENNSGTTCCRAVYS
						RVSAMTAPDKAFTPHRQRCRLP
						DATLARLIRPOIPAPSVGRIRRL
				1		RRIGNSANCLMRRLRVLSGHKF
				ļ		PHHPSMPNVSANGYRSFPFMAS
						YMTPFFGYLTAWSYCLCGCRW
				l		GSLKSPPLALCPVLVPGDGAVD
						TRIDAVALVALANLAAVRLYA
						KSSSAAGIINFVVLTAALLAVT
				1		AACTVATYALRTGKNROYRA
						MOSFRESSIDNEVDTFFKGTGA
		1				LGISVTVIAVTCKSRLVYPIPVY
						TGLSLPORPAVVKVNEGSMET
						KDLIVI/GAGINGAGIAADAAGR
	ļ					
						G/LSVLMLEAQDLACATSS/ASS KLIHGGLRYLEHYESACEALAE
				}		
						REVLLK/MAPHIAFPMRFRLPH
		l				RP/HLRPAWMIRIAENIS/DFVEP
		l				KLCLPRMGTRESVLPGVCRNY
						ESTITSVISVSPCERDMANYVPR
		l			•	HSAGYRRRFYPMTQPIKDAVEP
		<u> </u>				VDQDWKAGKEMAASRS
15924	46292	A	16015	1	5895	
15925	46293	Α	16016	170	371	
15926	46294	A	16017	216	477	
15927	46295	Α	16018	2	263	
15928	46296	A	16019	11	2862	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
15929	46297	Α	16020	1	1567	MKLNKAGFNVPESYSLLKMPP
				l		VGCLISALKKAEDRQEVILRLF
	i					NPAESATCDATVAFSREVISCSE
				l		TMMDEHITTEENQGSNLSGPFL
	Į.					RVRAGESIKFFNVLLADTPGLD
İ				l		DTMDKDVAHDSRSIQLAMLRD
						DEILTHPVFNRYHSETEMMRY
				l		MHSLERKDLALNQAMIPLGSCT
				l		MKLNAAAEMIPITWPEFAELHP
				l		FCPPEQAEGYQQMIAQLADWL
		1		l		VKLTGYDAVCMQPNSGAQGE
						YAGLLAIRHYHESRNEGHRDIC
				l		LIPASAHGTNPASAHMAGMQS
l				l		RKTAGICCVHLWAGFGKVAIIG
				ŀ		AGPAGLQASVTLTTQGYDVTIY
	i	ŀ		l		EKEAHPGGWLRNGIPQFRLPQS
1					l	VLDAEIARIEKMGVP\IKCTTEV
1						G\NTLTLEQVKAENRAVLVTVG
				l	1	LSSGSGLPLFEHSDVEIAVDFLQ
						RARQAQGDISIPQSALIIGGGDV
		1		l		AMDVASTLKVLGCQAVTCVAR
		1		l		EELDEFPASEKEFTSARELGVSII
						DGFTPVAVEGNKVTFKHGDLR
1						TAPFLGVADKRNKSAGNHP
15930	46298	A	16021	62	329	
15931	46299	В	16022	289	334	
15932	46300	Α	16023	1	2991	
15933	46301	Α	16024	418	3156	
15934	46302	Α	16025	1	2169	

2055

SEQ ID	SEQ ID NO:	Met	ISEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
1.502.5	146000	 	114004	lana	10.404	CORPA (FREE LAND LAND LAND LAND LAND LAND LAND LAND
15935	46303	A	16026	800	2406	SSRRMFPPAIRLLLICLPARRWK
		l		1		SAL/GEEQLQNLQHISQEMGISP
		l		l	1	SLFSTQPLWQIAMVLQATQAQ
		l				KLGLRAEYGIDYQLLQAAKQQ
	1	l		l		HKPVIELEGAENQIAMLLQLPD
	l	l		l		KGLALLDDTLTHWHTNARLLQ
		1		i		QMMSWWLECEGGERERIFPVS
		1		l		KRMGKKLDAVFGDNHTASLTE
		1				TEEYRKSQVHFCICDKRERSSG
						NYQSIQGCRSAAFFIPSNSTRTK
1	1	l				RGESMYRMDKLTTGAAYGAS
	l			ŀ		AGSILNGMLNAYSPEQWNAIG
						VLTTAAAGAEMNPTLRNKLVG
	1				İ	AIVGGSGAITIAAVMLGNADGL
	1	İ		l	ľ	EGRRYYAYQDVVGVWTVCDG
					ŀ	HTGTDIRRGHRYTDKECDNLL
	l					KADLRKAIIYISGYSGTNGIRDS
	l					LLFSSLWLIPVFLFPKRIKIIAAVI
1	l					GVAFSRFIASHLAVAFESDPEAE
l						IRQLNSRRVELERALSNHENDN
						QQQRIQFEQAKEGVTALNRILP
						RLNLLADDSLADRVDEIRERLD
				ì		EAQEAARFVQQFGNQLAKLEPI
						VSVLQSDPEQFEQLKEDYAYSQ
1						QMQRDARQPGVCP
15936	46304	Α	16027	2336	3105	HGNAIALIASQTWRDHPVDLLL
				l		SLLVQSLT/GRLRELNQSLQKEL
		l				ARNOHLAERLLETEESVRRDV
				l		ARELHDDIGQTITAIRTQAGIVQ
ŀ		l		l		RLAADNASVKQSGQLIEQLSLG
Į.		1				VYDAVRRLLGRLRPROLDDLT
						LEQAIRSLMREMELEGRGIVSH
		1				LEWRIDESALSENORVTLFRVC
						QEGLNNIVKHADASAVTLQGW
						QQDERLMLVIEDDGSGLPPGSG
				l		OOGFGLTGMRERVTALGGAFH
				1		ISCLHGTRVSVSLAORHV
15937	46305	c	16028	l	1374	
	10000				1071	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		ľ
		_				
15938	46306	A	16029	1298	2896	GAFTCCATAGGCMNIRRKNRL
						WIACAVLAGLALTIGLVLYALR
						SNIDLFYTPGEILYGKRETQQM
		l				PEVGQRLRVGGMVMPGSVQR
						DPNSLKVTFTIYDAEGSVDVSY
						EGILPDLFREGQGVVVQGELEK
		i				GNHILAKEVLAKHDENYTPPEV
						EKAMEANHRRPARPSIMMPEIG
						NGLLCLALGIALLLSVYPLWGV
						ARGDARMMASSRLFA WLLFMS
		l				VAGAFLVLVNAFVVNDFTVTY
						VASNSNTQLPVWYRVAATWG
					1	AHEGSLLLWVLLMSGWTFAVA
			Ì			IFSQRIPLDIVARVLAIMGMVSV
		i				GFLLFILFTSNPFSRTLHLAYPN
						RCQPINWRGRHLVWYLLLFVIS
1		1				LWLQLGLPDELSRFTPFCLALPI
		l				IALAWHYGWQGALIATLMNAI
		ł				ALIASOTWRDHPVDLLLSLLVQ
						SLTGLLLGAGIORLRELNOSLO
		l l				KELARNOHLAERVLETEOSVR
						RDVARELHYDIGOTITAIRTHA
		l				GIVORLAADNASVKOSGOLIEO
				ŀ		LSLGVYDAVRPLLGRLRPAOLD
		l		ŀ		DLT\LEQAIPLT
15939	46307	A	16030	15	277	KIGOKAPLGEARGGERT\SRTA
1		1		I		KNFGYEEMLSELEA/HRRG/WL
		1		İ		KRV*PRMKLPPEALVETFIAOLF
		1			1	DAPSQIMPLTSPIILLIAIEMGS
15940	46308	A	16031	1	2916	
15941	46309	A	16032	i	2862	
15942	46310	A	16033	i	3099	
13742	170310	111	110000	l <u>'</u>	12077	L

SEQ ID	ISEO ID NO:	Mar	SEQ ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
	l .			sequence		
15943	146211	-	16034	li -	1567	MKLNKAGFNVPESYSLLKMPP
15943	46311	A	10034	l'	1367	
	1	l	l			VGCLISALKKAEDRQEVILRLF
1		l				NPAESATCDATVAFSREVISCSE
		1				TMMDEHITTEENQGSNLSGPFL
		1				RVRAGESIKFFNVLLADTPGLDI
			İ			DTMDKDVAHDSRSIQLAMLRD
		1		l		DEILTHPVFNRYHSETEMMRY
		1		ľ		MHSLERKDLALNQAMIPLGSCT
						MKLNAAAEMIPITWPEFAELHP
		ļ		l		FCPPEQAEGYQQMIAQLADWL
		1				VKLTGYDAVCMQPNSGAQGE
		1		l		YAGLLAIRHYHESRNEGHRDIC
		1				LIPASAHGTNPASAHMAGMQS
		1	ļ			RKTAGICCVHLWAGFGKVAIIG
		1				AGPAGLQASVTLTTQGYDVTIY
		1				EKEAHPGGWLRNGIPQFRLPQS
		1		ł	\	VLDAEIARIEKMGVP\IKCTTEV
	1					G\ntltleqvkaenravlvtvg
		1		ŀ		LSSGSGLPLFEHSDVEIAVDFLQ
				ŀ		RARQAQGDISIPQSALIIGGGDV
		1				AMDVASTLKVLGCQAVTCVAR
1		l		1		EELDEFPASEKEFTSARELGVSII
		1		ļ		DGFTPVAVEGNKVTFKHGDLR
		1		1		TAPFLGVADKRNKSAGNHP
15944	46312	Α	16035	367	849	NNFCSTTSHHGQHHHTDHRCG
1		1				EQNVCHRDAICQTLFRTTKDDS
						NFICLIEAQSACHKDVNYNHDS
}		l		1		QKNAADHHDKQQ*SPANFMPD
		1				TINDGFAERCINHQQNGRLVSF
	1		l	[RNPFFIARHPVTNPRPGKMPQH
1		i				KRQQQLQHNFTDLLKAAPCTV
						NIHHQTNQQW
15945	46313	A	16036	1	1066	MHQRPRTRNAVEQHRFSRQLY
1						YALHAFEITIPPLRMRRGSIPAL
1		1		1	1	VNNKLRSLEKRFSTRLKIDDDA
		1	1	l		LARLVSCAWPGNDFELYSVIEN
		1	l	l		LALSSONGRIRVSDLPEHLFTEQ
1		1				ATDDVSATRLSTSLSFAEVEKE
		1		1		AIINAAQVTGAFPWWLGDVKL
1		1	l	l		TIPLTLGMVAAALTDLDDRLAG
				ł		RLRNLIITLFCFFIASASVELLFP
						WPWLFAIGLTLSTSGFILLGGLG
1		1		l		QRYATI\AFGA\LLNAIYTMLGT
1		1		l		SLYEHWYQQPMYLLAGAVWY
1		1		l		NVLTLIGHLLFPVRPLQDNLAR
		1		l		CYEQLARYLELKSRMFDPDIED
1		l	l	l		QSQAPLYDLALANGLLMATLN
		1	1	l		QTKLSLLTRLRGDRGQRGTRRT
		1		l		GIL
15946	46314	A	16037		951	OIL .
15946	46314	A	16037	576	669	
13947	40313	I'M	10038	270	007	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15948	46316	Α	16039	jı .	1047	
15949	46317	Α	16040	2	425	
15950	46318	Α	16041	148	1052	RARRGVSVKAKKQETAATMK
						DVALKAKVSTATVSRALMNPD
						KVSQATRNRVEKAAREVGYFP
						QPMGRNVKRNESRTILVIVPDIC
						DPFFSEIIRGIEVTAANHGYLVLI
		l				GDCALQIQQEKPFIDLIITKQIDG
						MLLLGSRLPFDASIEEQRNLPP
						MVMANEFAPELELPTVHIDNLT
						AAFDAVNYLYEQGHKRIGCIA
						GPEEMT\LCHYRLQGYVQALRR
						CGIMFCRFTAEGVQEIDYKDIA
						TLKNYITESGGVTELIHFFIAEY
		l				SDNQRANAGGGVEDEDIEVLES
		l				LRLNENHAIKERCRS
15951	46319	Α	16042	3	741	
15952	46320	A	16043	237	1542	FHRLWTGRFLPSPPSRRGTDNA
		l				AGRTKMPPSPVIF/WFALAQFV
	l	1			Į.	GVKHSRSIPFAAVRSHKARIRSI
		1				SRSDVATSILPQLS*RMPF*SQK
	ł	l				ALVARFPARQNCAFRLPGV*/L
	ļ					DPGMDHAAVVARLVTGRGGIV
		1				WFGRDFYRSVDDIHFGAWGDL
]				AKECFDIVIPQTNTARTYPHAN
		1				AEIGVRTMKQINAAIGRQTNRM
	l	1				MAHWVIRSSRHYRRQLNSLCFI
		l				TGADVCRWFPQHIVPGRDSVA
		1				HVVVQHVITYRLLRGLLHFAG
						DGGVNAITVFIGTFAITVIHLLA
		1		l		HHFAQVRRRESNLRRVIVSVNR
						LSACLIELLLGDIPFIQHPRQHD
	ŀ	l			ł	VTPRDGTIHRVERVEGGRRFRQ
		1				PGYHRHFVQRQLVDRFSEIDLC
		l				RSTDTIGAVTKVNLVQIKLEDL
		1				VFRQQLFNANREEDFLNLTHQR
		_				AFWTEEEVSRQLLGNGTCPL
15953	46321	Α	16044	1	1059	
15954	46322	A	16045	104	349	RSAASPWKTRKALVCCSAVVIS
				l		PALKISMIVTISVQAHG*LRKGS
	1	1	İ	l		GVKAALSWWKFQPTMKPTITS
l		_				SLTGRRISCRSRVKR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown,
15955	46323	A	16046	1056	2373	GPRLVORSDHRPHDANYROYL AGISVHODPLHGLAENTIPPDL YEASAMDGAGPFQNFFKITLPL LIKPLTPLMIASFAFNNFVLI LUNYTYRIAFEGGGGDFGLA AAIATLIFLLIRPRATTALGOTDL LUNYTYRIAFEGGGGDFGLA AAIATLIFLIRIPRATTALGOTTDL LUNYTYRIAFEGGGGDFGLA KILMESSRWARANRILKAIN TILSLITITOSCETYLIQNTRPELI TOTFREFFDTPYETAQDVHKQL KRLRRVIAWTGERETPVTIYSW VAAATRYQLLKRGVISNTKAIN TEEEILQGEPEVKVESAERHHA MVNFWRTTLSCLIGTLFWLWT GGGRPVAVGVFYGGCYCTRDG LTRAPPPNAAPTAARYGPRTTP RTLRKGITDVRCEKRHIRGRQA RNAGYGTVFSSLDRLRHPALR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \≔possible nucleotide insertion)
15956	46324	A	16047		2581	MVLMGLAQNIWOFLILRALLG LLGGFVPNANALIATQVPRNKS GWALGTLSTGGVSGALLGPMA GGLLADSYGLRPVFFTTASVLIL GFVTLECIREKFOPVSKEML HMREVVTSLKNPKLVLSLEVTT LIIQVATGSIAPILTLYVRELAG NYSNVAFISGMIASVPGVAALL SAPRLGKLGDRIGPRKILITALI SAPRLGKLGDRIGPRKILITALI SAPRLGKLGDRIGPRKILITALI NKSNVAFISGMIASVPGVAALL SAPRLGKLGDRIGPRKILITALI NKSNVAFISGMIASVPGVAALL SAPRLGKLGDRIGPRKILITALI THOMINAY WTPDOLPEPKEM NKYTITTSRDEDKLHAPDNAW VQQTRSTGDVKQSNLIRQPDG TIAFVVDFTCAGMKKLPGDTY TAQTSIGDNGEIVESTVRYNPV TKGWRLVMRKVKADAKTITE KGWRLVMRKVKADAKTITE KGWRLVMRKVKADAKTITE KGWYLYGPLDSIWAFGLGANYVKQ RDWRSAKDMMKFTDYSVKTG HLTATWTFSFAQDVLKASVG QVLAGDKGTLEIAKREDSGV VVGGYATITNYSKEEVGEODFI KGYVYSVPLDSSGPTRSRAA IGWTPLTRDGGQLGPGYTDLL VNYTYRIAFEGGGQLGPGYTDLL VNYTYRIAFEGGGGQFGLAA IATLIFLLVAIMPPLLMVVAIS LRAGFSVEROAGRITPPFPVLL WLWNSVKVAGISAIGIVALSTT
		ļ.		ļ	2/4	CAYAFARMRFPGKATLLKGML
15957 15958	46325 46326	A	16048 16049	869	765 1012	MAISFMMSCFVT*VAANSAILL SMSWRA*SNSNGP*LANSILLS AAC

SEQ ID	ISEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
	<u> </u>	<u> </u>			1	New contame and the contame
15959	46327	A	16050	1	1854	MTLSCNSTDFEKTVTSVPALPD
1		1			ĺ	VVSDSDMSLNALSVTLADNVQ
		1			!	LSFRRDSYSGQPSVIYTTPQRPT
			1			AGSEDAPVWPEAFTEDVRHNR
	1	ı				PPYPVIPRATRPACLARERMAM
	1	i				YMRPQIKTIGKAPAEEALSNPR
		1				MQGPQLSSASTIPVTPFHWLFG
		1				SHAAGHFSQHAHRTALTGHFS
į.		1				HLATTLKGLALEGNAIFTITRFT
	i	1				RPYGHKNTTPHSPLCSAPHSPIF
		İ				NSRRKVYGQHGPQWWAFIHSH
		1				RPGGSSHSSILHCFALVCPGLVV
		1				QRDTPHPIIRPALCPPIAIAFSLSL
		1				WLGNPSFPSGTQHLQGPKTFQC
1						OASPSSKVCPKRLICTVIPOORV
1		1	1			ASHHHFHGHNRYYQLGLPVQH
l .		1				CTLOGVIRATHFIPRTARNOISH
		ì	1			VSWGAGPAHSDSQGKQDLE/H
		1				EPIKPLDNSEKASKV/SOALETV
						TTTAEKVQRQ/PVIAHLIRATER
		1				FNDR/LGNQFGGFVLAS/HPMLL
1						QDIFDKILQNIS/DPTLAATLKNT
			1			INTAVHSHLVPAGADGIFLLLRI
						FDAVCAAWIATAEYKDYPGMP
]			i		GKTHLKIRLMRRERLIGIRFYSG
1	1					LLPGTKAPYSRTLAFKRVLHTA
1						
1						LSLHLGLGKTCTTGFELCSLTLI EYASNAFTYWPDWHINRIKNK
						N PROPERTY WEDWHINKIKNK
15960	46328	A	16051	249	724	EKVKKTQRKEIENVTNITGVRQ
13900	40328	^	10051	-77	1/27	IELWRRDDLQHPRLDEVAEEVE
		1				VALVYNGISHVVMMASPKDLE
1						YFALGFSLSEGD/IIESPRDIFGM
						DVGVPCACNGLEERRRALDG/R
						TVAGCGVCGVEQ/LLNDIGKPV
		i	1	1		QPL\PFTQT/FDLNKLDDALRHL
1		1		1		NDFOPVGN
15061	46329	A	16052	71	408	RYLSTASQTNSIQIGAGLSRSRA
15961	40329	I ^A	10052	 ''	1400	
		i				FTTFGSKGILLSHTSVFIPRDRCT
1		1				SFSMMPTCSASLTSHCPFQMTS
1	1	1	1	1		LLSSHHP/FKKKKFGT\KTGHCL
L	ļ	1				CPSLNCLPCCFSQIMVIKQVIL
15962	46330	Α	16053	668	905	CLHVLHHLTSHCPFQMTKPYST
	1	1				IHFLPRMTSLLSSHHP/FKKKKF
		1	1	1		GT\KTGHCLCPSLNCLPCCFSQI
L	L	\bot				MVIKQVILEKKKK
15963	46331	Α	16054	834	1057	ARLLFMDLLDERTLVAVERPL\
		1				VTSLLSSHHP/FKKKKFGT\KTG
		1				HCLCPSLNCLPCCFSQIMVIKQV
		١.				ILEKKKKTK
15964	46332	Α	16055	1	1011	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15965	46333	Α	16056	191	631	NLLRGNKMATQSSPVITDMKVI
		1			\	PVAGHDSMLLNIGGAHNAYFT
						RNIVVLTDNAGHTGIGETPGGD
		l				VIYQTLVDA\IPMVLGQEVARL
						NKMVPOVHKGNOAADFDTFG
						KGAWTFELRVEAGAGLEARLL
		l		ļ		DLLGKALNVSGCELLGHGS
15966	46334	Α	16057	2	335	GARQQOKFVLIAGGFP*QLL*T
		1				NAVIGREFRNVOECIGLHTAGF
		ı				ANAVTKHFGKCAIALHNMAFR
						IEDNHRSVEMVNGNNOLSGLFF
		1			1	LRHFYAGIPYOROVINFPLGAH
15967	46335	Α	16058	736	910	RVSTPRATGTVPQRDYPLSVVA
						TG*PDSFHTYLFVRTY*K*V*VL
						AYLNTRQNDGDFP
15968	46336	В	16059	17	1699	
15969	46337	Α	16060	3	440	
15970	46338	A	16061	1	564	
15971	46339	A	16062	1	1062	
15972	46340	Α	16063	203	425	LNNWIISWLLHMMCLMKVAN
						CFRS*DRRELAVSAFFVLL*AM
		l				RLRRSSSGFLANLAHLQKMKL
			L			NSSTYVPPTKH
15973	46341	Α	16064	402	713	RGVAISLGQSKSHGYIHFQVAQ
						REPEILG\DSIKAVRWCSLHGRL
		l	ļ.			M***M*RDLKQWFSAPEA*VSL
		l				GTLCVIGQKAVGSLALMSWEP
						TPRPMGLLSTSRTLYS
15974	46342	Α	16065	944	1305	LPLWFISALRLAGIAVWCVLST
		l		l	1	GFEQIRMIKAFGLEDRQSAL/FA
				l		ADAEDTGKKNMRVAR/IDARF
	1			l		DPTIGWPSSLGLLFSNRTPWVPF
				l		PFFFQVPQ\LHTFIRGILIWKTFP
		ı	1	l		FWIFPFPASSW
15975	46343	Α	16066	1	756	

15976	SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
15976							*=Stop codon, /=possible nucleotide
IS976 46344 A 16067 S0 III91 CLTIAVFFPAPQCPEMFSIP ARDGERAASMIQDYIFCFFF MKLKPERMISHERAQTSD DCRRHLPIGILMAGGMFAW KHQGQTFIERPRISLTCMLP ATTH-WIFTRIGQVLYTCLLI ECTV-NEV-QVJATVVSVNPP PDIVAMIGASAALSLSGIPF GAARVGVINDQVJANFOQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVFGHI QVDSRGSPIRGNRTCDNYT LIDYVNDMIENVSEEAFCV IMRISDHISGUVYCLSLFA LVLVVIAVSVVNPCLLEVTI AASINSQPNLNISDTWGIGA GAYTCSCLDPHEMMSSGF SHSCVEK LILFMST/ALLDAVVKKNRT PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI VYAFLGVPANLLMRKLGAI GTTTLLWGFLSAAMAWAI AKFLIVATLLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIEGILAV GVFTFFWLDDTPEQAFFLS KTLLINQLASEQQKVTSRI ALRNGRVWQLAITYLTIQV GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALLGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALGTKVGF SVVTAIPWVAALGGTWLPITUTQU GLIFFLDTQVAALGTVLDTUTQU GLIFFLDTQVAALGTVLDTUTQU GLIFFLDTQVAALGTVLDTUTQU GLIFFLDTQVAALGTVLDTUTQUTQUTQUTQUTQUTQUTQUTQUTQUTQUTQUTQUTQ		sequence	1	09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
ARDGERAASMIQDYIFCFFS MKLKPERMISHERAQTSDI DCRRHLPIGILMAGGMFAV KHQQGTIFERRISLTCMLP ATPLWLFTRIGQVLYTCLLI ECFVNEVQVIATVVSVNPC PDIVAMIGASAALSLSGIPFI GAARVGVINDQVJNPTQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVYGFH QVDSRGSPIKGNNTCDNYI LIDYVNDMIENVSEEAFCVI IMRRISDHIISCIVYCCLSLFA LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSG SHSCVEK ILILMSTIALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAVALGAGII VYAFLGVANLLMARLGAI IGTTTLLWGLSAAMAWAI AKFLIVRTLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIEGILAV GVFTFFWLDDTPEQAFLSI KTLLINQLASEEQKVTSRI ALRNGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTVLIPU DKSAAKRYTRPHRECGYSE IIPPKKVRNNAPHSGPGPFICI RVRVRNLGIEVYKHDSPPP			l		sequence		
ARDGERAASMIQDYIFCFFS MKLKPERMISHERAQTSDI DCRRHLPIGILMAGGMFAV KHQQOTTERERRISLTCMLP ATPLWLFTRIGQVLYTCLLI ECFVNEVQVIATVVSVNPC PDIVAMIGASAALSLSGIPFI GAARVGVINDQVJVNPTQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVFGH QVDSROSPIKGNNTCDNYT LIDYVNDMIENVSEEAFCVI IMRRISDHISICTVYCCLSLF LVLVVIAVSVVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSG SHSCVEK TYQIDTGISNEAVALGAGII TYQIDTGISNEAVALGAGII TYQIDTGISNEAVALGAGII TYQIDTGISNEAVALGAGII TYQIDTGISNEAVALGAGII GTTTLLWGFLSAAMAWAI AKFLLYRTLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIEGILAV GVFTFFWLDDTPEQAFLSI KTLLINQLASEEQGKVTSRI ALRNGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTVAIPL DKSAAKRYTRPHRECGYSE IIPPKKVRNNAPHSGPGPICI RVRNRNLGIEVYKHDSPP	15076	46244		16067	160	1101	CLTIA VEEDA DOCDEMENDOUT
MKLKPERMISHERAOTSD DCRRHLPIGILMAGGMFAV KHQQQTTELERRISLTCMLP ATFLWLFTRIGQVLVTCLLL ECFVNEZVQVLATVSVNPC PDIVAMIGASAALSLSGIPF GAARVGYINDQYVLNPTQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVCGHI QVDSRGSPIIGNRTCDNYI LIDYVNDMIENVSEEAFCV. IMRHSDHIISCIVVRCLSLFA LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSG SHSCVEK LILFNST/ALLDAVVKNRN PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGII VYAFLGVPANLJMRKLGAI IGTTTLLWGFLSAAMAWAL AKFLIVRTLRAAEAGFFC YLTSQWFQRYRASIMGLF GAPLALTLGSPLSGALLEMM MGHPGWFWMFVIGGLAV GVFTFWLDDTPEQAFFLSI KTLLINQLASEQQVKTTSM ALRNGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTVAIPL DKSAAKRYTRPHRECGYFS IIPPKKVNNAPHSGCIPGI RVRVRNLAIGVVKNNEPISPCIPC RVRVRNLAIGVVKNSPSPPRECGYFS IIPPKKVNNAPHSGCIPGTE RVRVRNLAIGVVKNSPSPPRECGYFS IIPPKKVNNAPHSGCIPGTE RVRVRNLAIGVVKHNSPP	13976	46344	l^	10007	30	1191	
DCRRHLPIGILMAGGMFAY KHQQTFIERPRISTCMLP ATFLWLFTRIGQVLVTCLLI ECFVNEZVQVIATVVSVNPP PDIVAMIGASAALSLSGIPFI GAARVGVINDQYVLNPTQI KESKLDLVVAGTEAAVLM EAQULSEDQMLGAVVFGHI QVDSRGSPIKGNRTCDNYTI LIDYVNDMIENVSEEAFCVI IMRRISDHIISCIVVRCLSLF, LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHRMMSSGF SHSCVEK I5977 46345 A 16068 401 1518 LILFMSTALLDAVVKNRN PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAVALGAGI UYAFLGVPANLLMRKLGAI VYAFLGVPANLLMRKLGAI GTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFPFP YLTSQWFPORNASIMGLF GAPLALTLGSPLSGALLEM MGHPOWFWMFVIEGLLAV GVFTFFWLDDTFEQARFLSI KTLLINQLASEEQOKVTSRI ALRRIGWWQLAIJVLTIOV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTWLPI DKSAAKRYTRPHRECGYFS HPPKKVRNAPHSGCPGIC RVRKPNLGIEVKHOSPP							
KHQGOTTERRRISLTCMLP ATFLWLFTRIGQVLVTCLLI ECTYNEVQVIATVVSVNPC PDIVAMIGASAALSLSGIPF GAARVGYINDQVJVAPTQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVFGHI QVDSRGSPIKGNRTCDNYT LIDYVNDMIENVSEEAFCV IMRISDHISICIVYCLSLFA LVLVVIAVSVVNPCLLEVTI AASINSQPILNISDTWGIGA GAYTCSCLADPHEMMSSGF SHSCVEK LILFMSTALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI UYAFLGVPANLLMRKLGAI GTTTLLWGFLSAAMAWAI AKFLIVRTLLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMYEIGLLAV GVFTFFWLDDTPEQAFLSI KTLLINQLASEQGVTSRI ALRNGRVWGLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTWLPILD DKSAAKRYTRPHRSCGYFS IIPPKKVNNAPHSGPGPICI RVRVRNLGIEVYKHDSPP	l		l				
ATELWIFTRIGOVLYTCLLI ECFVNEVQVIATVVSVNP DDIVAMIGASAALSLSGIPFI GAARVGYINDQYVLNPTQI KESKIDLVVAGTEAVLM EAQLLSEDQMLGAVVCFHI QVDSRGSPIKGNRTCDNYV LIDYVNDMIENVSEEAFCVI IMRRISDIHISCIVVRCLSLF, LVLVVIAVSVVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPIKMMSSGF SHSCVEK IS977 46345 A 16068 401 1518 LILFMSTALLDAVVKKINA ILFMSTALLDAVVKKINA VAFIGVPANLAMRALGAI IGTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPP YLTSQWFPQNRVASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIGGLAV GVFTFFWLDDTPEQARFLSI KTLLINQLASEQQVCTNSI KTLLINQLASEQQVCTNSI ALRINGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKYG SVVTAIPWVAALGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNAPHISGPGIC RVRKVRNADIEGGYVKHDSPP			l				
ECFVNEZ/QVIATVVSNING BCFVNEZ/QVIATVVSNING PDIVAMIGASALSLSGIPF GAARVGYINDQYVLNPTQI KESKLDLVVAGTEAVLM EQULSEDQMLGAVVFCH QVDSRGSPIKGNRTCDNYI LIDYVNDMIENVSEEAFCVI IMRISDHIISCIVYCLGLEF LVLVVIAVSVVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPIKMMSSGF SHSCVEK SHSCVEK LILPNST/ALLDAVVKINR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI VYAFLGVPANLLMRRLGAI IGTTTLLWGFLSAAMAWAI AKFLIYRTLRAARAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSFLSGALEM MGHPGWFWMFVIEGLLAV GVFTFFWLDDTPEQAFFLS KTLLINQLASEEQKVTSRI ALRNGRVWGLAITYLTIQV GLIFFLFTGVAALLGTKVGF SVVTAIPWVAALGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGFICI RVRVRPNLGIEVYKHDSPP							
PDIVAMIGASAALSILSIGIPE GARRYGYINDQYVLNPTQ KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVYCHI QVDSRGSPIIGNRTCDNYI LIDYVNDMIENVSEEAFCV IMRHSDHIISCIVVRCLSLFA LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIAPPHKMMSSG SHSCVEK LILFMSTJALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGII UYAFLGYANLAMIARKLGAI GTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPC YLTSQWFPQRYRASIMGLF GAPLALTLGSPLSGALLEMI MGHPGWFWMFVIGGLAV GVFTFWLDDTPEQARFLSI KTLLINQLASEEQQKVTSI KTLLINQLASEEQGKTSI KTLLINQLASEEQGKTSI ALRNGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTVAIPU DKSAAKRYTRPHRECGYFS IIPPKRVRNAPHSGPGIPE RYRVRNLGIEVYKHDSPP RYRVRNAPHSGFGOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIPPKRVRNAPHSGFOYFS IIP			l				
GAARVGYINDQYVLNPTQI KESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVFGHI QVDSRGSPIKGNRTCDNYYI LIDYVJADMENSVESEAFCVI IMRISDHIISCIVVRCLSLFA LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHIKMMSSGE SHSCVEK LILFMST/ALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI VYAFLGVPANILMRKLGAI IGTTILLWGFLSAAMAWAI AKFLIVRTLRAAEAGFPFP YLTSQWFPQNTRASIMGLF GAPLALTLGSPLSGALLEM MGHPQWFWMFVIEGLLAV GVFTFWLDDTPEQARFLSI KTLLINQLASEEQQKVTSRI ALRNGRWQGLAHJVTIQV GLIFFLFTQVAALLGTKVG SVVTAFPWVAALGTWLPI DKSAAKRYTRPHRECGYFS HIPPKKVRNNAPHSGOPGIC RVRWRNLGIEVYKHDSPP	1		1				
RESKLDLVVAGTEAAVLM EAQLLSEDQMLGAVVFGH QVDSRGSPIKGNRTCDNYI LIDYVNDMIENVSEEAFCVI IMRISDHIISCIVYCLGLEF LVLVVIAVSVVNPCLLELFI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSG SHSCVEK IS977 46345 A 16068 401 1518 LILFMST/ALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAVALGAGII VYAFLGENBAVALGAGII VYAFLGENBAVALGAGII GTITLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWHYEIGLAV GVFTFFWLDDTPEQAFLSI KTLLINQLASEQQKVTSRI ALRNGRVWGLAITYLTIQV GLIFFLFTQVAALLGTKVGF SVVTAIPWVAALGTKVIG SVTAIPWVAALGTKVIG SVTAIPWVAALGTKVIG FURNRAFINGGIPLIC RYRVRPNLGIEVYKHDSPP	ŀ		l				
EAQLLSEDOMLGAVVEGH QVDSRGSPIKGNRTCDNYY LIDYVNDMIENNSEEAFCVI IMRRISDHIISCUVYRCLSLF LVLVVIAVSVVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPIKMMSSGF SHSCVEK LILFMSTJALLDAVVKKMSK TYQIDTGLSNEAYALGAGI TYQIDTGLSNEAYALG			ĺ				
QVĎSRGSPIKGNRTCDNYT LIDYVNDMENVSEEAFCVI IMRHSDHISCIVVRCLSLFA LVLVVIAVSVVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPIKMMSSGP SHSCVEK I5977 46345 A 16068 401 1518 LILFMST/ALLDAVVKKNR PYMLAL VVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI VYAFLGVPANLLMRKLGAI IGTTTLL WGFLSAAMA WAI AKFLIVRTLLRAAEAGFPFC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHBGWFWMFVUEGLLAV GVFTFFWLDDTPEQAFFLS KTLLINQLASEEGQK VTSRI ALRNGRVWQLAIIYLTIQV GLIFFLFTQVAALLGTKVGF SVVTAIPWVAALGTWLIPL DKSAAKRYTRPHREGCYSE IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVVKHDSPP							
LIDYYNDMIENVSECAFCV IMRHSDHIISCIVVRCLSLFA LVLVVIAVSVVNPCLLEVTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHIKMISGG SHSCVEK IS977 46345 A 16068 401 1518 LILFMST/ALLDAVVKKNN PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGII VYAFLGYPANLLMRKLGAI IGTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPC YLTSQWFPQRYRASIMGLF GAPLALTLGSPLSGALLEMI MGHPGWFWMYEIGLAV GVFTFEWLDDTPEQAFFLSI KTLLINQLASEEQOKVTSI ALRNGRVWQLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTWLPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGIPE RVRVRNLGIEVYKHDSPP			ı				
IMRHSDIHISCIVVRCLSEA LVLVVIAVSVNPCLLEVT AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSG SHSCVEK 15977 46345 A 16068 401 1518 LILFMST/ALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGII VYAFLGVPANLLMRKLGAI IGTTILLWGFLSAAMAWAI AKFLIVRTLRAARAGFFP YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEMI MGHPGWFWMFVIEGLLAV GVFTFWLDDTPEQARFLSI KTLILNQLASEEGOK VTSRI ALRNGRVWQLAIIYLTIQV, GLIFFLFTQVAALLGTKVG SVVTAFWVAALGTWLIPU DKSAAKRYTRPHREGCYFS IHPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP			l				
LVLVVIAVSVVNPCLECTI AASIVSQPNLNISDTWGIGA GAYTCSCIADPHKMMSSGG SHSCVEK I5977 46345 A 16068 401 1518 LILFMSTALLDAVVKKNR PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGII VYAFLGVPANLLMRRLGAI IGTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVLGGLAV GVFTFFWLDDTPEQAFFLS KTLLINQLASEEQKVTSRI ALRNGRVWGLAITYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTWLPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP			l				
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GAYTCSCIADPHKMMSSGF SHSCVEK 15977 46345 A 16068 401 1518 LILIFMST/ALLDAVVKKNRN PYMLALYVLAFLDRSNIGL TYQIDTGLSNEAYALGAGI VYAFLGVFANLLMRKLGAI IGTTTLLWGFLSAAMAWAI AKFLIVRTLLRAAEAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIEGLLAV GVFTFFWLDDTPEQAFFLSI KTLLINQLASEEGOK VTSRI ALRNGRVWQLAIIYLTIQV, GLIFFLFTQVAALLGTKVGF SVVTAIPWVAALGTWLIPL DKSAAKRYTRPHREGCYFS IHPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							
SHSCVEK							AASIVSQPNLNISDTWGIGAILY
IS977 46345 A 16068 401 IS18 LILFMST/ALLDAVVKKNRN PYMLALYVLAFLORSND TYQUDTGLSNEAYALGAGII VYAFLGYPANLLMRKLGAI IGTTILLWGFLSAAMA WAI AKFLIVRTLLRAAEAGFFPC YLTSQWFPQNRVASIMGLF GAPLALTLGSPLSGALLEMI MGHPGWFWMFVIEGLLAV GVFTFFWLDDTPEQARFLSI KTLLINQLASEEGOKVTSRI ALRNGRVWQLAIIYLTIQV, GLIFFLFTQVAALLGTKVG SVVTAFPWVAALGTKVG SVVTAFPWVAALGTKVG SVTAFPWVARLGTWLFIP DKSAAKRYTRPHRECGYFS IHPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							GAYTCSCIADPHKMMSSGPMY
PYMLALYVLAFLDRSNIGL TYQIDTGISNEAYALGAGII VYAFLGUSNEAYALGAGII VYAFLGUSNEAVALGAGII VYAFLGUSNEAVALGAGII VYAFLGUSNEAVALGAGII IGTTILLWGLSAAMAWAI AKFLIVRTLLRAARAGFFPC YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVLGGLAV GVFTFFWLDDTPEQAFFLSI KTLLINQLASEEQKVTSRI ALRNGRVWQLAIIYLTIQV GLIFFLFTQVAALLGTKVGF SVVTAIPWVAALGTKVGF SVVTAIPWVAALGTKVGF FUNGAGIIFLFTGVAGIIGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							
TYQIDTGLSNEAYALGAGI VYAFLGVPANLLMRKLGAI IGTTTLLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFPP YLTSQWFPORTNASIMGLF GAPLALTLGSPLSGALLEMI MGHPOWFWMFVIGGLAV GVFTFFWLDDTPEQARFLSI KTLLINQLASEEQOKVTSRII ALRNGRWQULAHIYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTKVG FVFTFWLDTFEQARFLSI HPFKTVRNAFHSGFØFIC RVRVRNLGIEVYKHOSPP	15977	46345	Α	16068	401	1518	LILFMST/ALLDAVVKKNRVRLI
VYÄFLGYPANLLMRKLGAI IGTTILLWGLSAAMAWAI AKFLIVETLERAAGGFPE YLTSQWFPORNRASIMGLE GAPLALTLGSPLSGALLEMI MGHPGWFWMFVIEGLLAV GVFTFFWLDDTPEQARFLSI KTLLINQLASEEGOK VTSRI ALRNGRVWQLAIIYLTIQV, GLIFFLPTQVAALLGTKVG SVVTAIPWVAALGTKVG FYVTAIPWVARLGTWLPI DKSAAKRYTRPHREGGYPS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							PYMLALYVLAFLDRSNIGLAKQ
IGTTILLWGFLSAAMAWAI AKFLIVRTLRAAEAGFFFC YLTSQWFPQRYRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVIGGLAV GVFTFWLDDTPEQARFLSI KTLLINQLASEQQKVTDIST ALRNGRVWQLAIIYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTKVG FVFTFWLDTPEQCFS IIPPKRVRNAFHSGFQFS IIPFKRVRNAFHSGFGTE IPFKRVRNAFHSGFGTE RVRVRPNLGIEVYKHDSPP							TYQIDTGLSNEAYALGAGIFFV
AKFLIVRTLLRAAEAGFFPC YLTSQWFPQRYRASIMGLE GAPLALTLGSPLSGALLEM MGHPOWFWMFVIEGLLAV GVFTFWLDDTPEQARFLSI KTLLINQLASEEQOKVTSRI ALRNGRVWQLAIIYLTIQV GLIFFLPTQVAALLGTKVGF SVVTAIPWVAALFGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP			1				VYAFLGVPANLLMRKLGARTW
YLTSQWFPQRNRASIMGLF GAPLALTLGSPLSGALLEM MGHPGWFWMFVLGGLAV GVFTFFWLDDTPEQAFFLS KTLLINQLASEEQK VTSRI ALRNGRVWQLAHYLTIQV, GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTKVGF FOXATAPWAALGTKVGF HPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP			l				IGTTTLLWGFLSAAMAWADTE
GAPLÄLTLÖSPLSGALLEM MGHPOWFWMFVIEGLLAV GVFTFFWLDDTPEQARFLSI KTLLINQLASEEQQKVTMIN ALRMGRVWQLAIFYLTIQV GLIFFLFTQVAALLGTKVG SVVTAIPWVAALGTKVG DKSAAKRYTRPHRECGYFS HIPPKKVRNNAPHSGPÖPICL RVRVRNLGIEVYKHDSPP							AKFLIVRTLLRAAEAGFFPGMI
MGHPGWFWMFVIEGLLAV GVFTFWLDDTPEQARFLS; KTLLINQLASEEGORVTSRI ALRNGRVWQLAIIYLTIQV, GLIFFLPTQVAALLGTKVG SVVTAIPWVAALIGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP			1				YLTSQWFPQRNRASIMGLFYM
GVFTFEWLDDTPEQAEFLS KTLLINQLASEQOKVTSRI ALRNGRVWQLAIIYLTIQV. GLIFFLDTQVAALLGTRVG SVVTAIPWVAALGTRVG DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGFQDIC RVRVRPNLGIEVYKHDSPP							GAPLALTLGSPLSGALLEMHGF
KTLLINQLASEQQKVTSRI ALRNGRVWQLAITYLTIQV/ GLIFFLFTQVAALLGTKVG SVVTAIPWVAALFGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							MGHPGWFWMFVIEGLLAVGA
ALRAGRVWQLAITV.TIQV. GLIFFLPTQVAALLGTKVG SVVTAIPWVAALFGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPP							GVFTFFWLDDTPEQARFLSKQE
GLIFFLPTQVAALLGTKVGF SVVTAIPWVAALFGTWLIPI DKSAAKRYTRPHRECGYFS HIPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPPI							KTLLINQLASEEQQKVTSRLSD
SVVTAIPWVAALFGTWLIPI DKSAAKRYTRPHRECGYFS IIPPKKVRNNAPISGOPFICI RVRVRPNLGIEVYKHDSPP							ALRNGRVWQLAIIYLTIQVAVY
DKSAAKTYRPHRECGYFS HIPPKKVRNNAPHSGPDIC RVRVRPNLGIEVYKHDSPP	i		l		1		GLIFFLPTQVAALLGTKVGFTA
HPPKKVRNNAPHSGPGPICI RVRVRPNLGIEVYKHDSPPI	l		ĺ		1		SVVTAIPWVAALFGTWLIPRYS
RVRVRPNLGIEVYKHDSPP			l		1		DKSAAKRYTRPHRECGYFSAPY
			l		1		HPPKKVRNNAPHSGPGPICLIRL
	l						RVRVRPNLGIEVYKHDSPPNNA
15978 46346 A 16069 283 822	15978	46346	A	16069	283	822	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
15979	46347	IA	16070	II	1400	MTDLIQRPRRLRKSPALRAMFE
13979	46347	A	10070	1	1400	ETTLSLNDLVLPIFVEEEIDDYK
		l				
			1			AVEAMPGVMRIPEKHLAREIER
		l			l	IANAGIRSVMTFGISHHTDETGS
					1	DAWREDGLVARMSRICKQTVP
		ļ				EMIVMSDTCFCEYTSHGHCGV
					l	LCEHGVDNDATLENLGKQAVV
					ŀ	AAAAGADFIAPSGAMDGQYDV
						FVRNENYWGEKPAIKKITFNVI
						PDPTTRAVAFETGDIDLLYGNE
		l				GLLPLDTFARFSQNPAYHTQLS
		l		Ì	İ	QPIETVMLALNTAKAPTNELAV
		1			ł	REALNYAVNKKSLIDNALYGT
		l				QQVADTLFAPSVPYANLGLKPS
		l				QYDPQKAKALLEKAGWTLPAG
		l			<u> </u>	KDIREKNGQPLRIELSFIGTDAL
1		l				SKSMAEIIQADMRQIGADVSLI
			l		i	GEEESSIYARQRDGRFGMIFHR
		l				TWGAPYDQRPSQSKRVRQR*F
		l				QAQQD*MKKPTQSKWNRATR
		l	1			VRGGREILTKFVLTAPLKWLKR
		1			İ	SNSRRVSFAG
15980	46348	Α	16071	592	1427	IPIGVTPCFSTLRRTLFALLACAS
		1				FIVHAAAPDEITT\AWPVNVGPL
		1				NPHLYTLNQMFAQSMVYEP\LV
		i i				KYQADGSVIPWLAKSWTHSED
1		1				GKTWTFTLRDDVKFSNGEPV\D
		l				AEAAAENFRAVLDNRQRHAW
		l	ļ			LELANQIVDVKALSKTELQITL
		l				KSAYYPFLQELALPRPFRFIAPS
		l				QFKNHETMNGIKAPIGT\QPWN
		l	1			LQESKLNQYHGFGRNENYWGE
		l				KPAIKKITFNVIPDPTTRAVAFE
		l				TGDIDLLYGNORLLPLDTFARF
		1				SQNPVYHTQLSQIWNQ
15981	46349	Α	16072	I	1071	
15982	46350	Α	16073	585	1072	RPARDVPLPIRGLQPSLRGPQQL
		l				PEKPAVEVRLDKWLWAARFYK
		1		1		TRALA\REMIEGGKVHYNGQRS
	1	1				KPSKIVELNATLTLRQGNDERT
						VIVKAITEORRPASEAALLYEET
						AESVEKREK\MSPRGPPFVGAP
1		1				KNPRGFLKLNALTMPHPDRRP
1						DKKERRDLLR
15983	46351	A	16074	36	356	ASHTLARKSC*S/PSLLYQENQA
1		Γ.		F-	I	HNQMPASELKASEIPFHPSIKTQ
						DPKAEEKSPRSKR*L*QRQVLP
1		l	l			DPAADLSVKPPGPLMLPLCYLS
						GPADITVLKASQDGLIIA
		Ц_	L	L		OI ADII YERASQUOLIIA

15984 46352 A 16075 I 2111 MNTPLQSAHTHLT	
SDSPFPAASTLSLP KLGAGTMPPAAC PPAAYPHRNKVRA PPAAYPHRNKVRA PVREYEAGDEASG KSVQNARADRGM VRGGENTVVVFAA QIVLRDIAREMY TQMDAKKPRKCDI RKKQRFTSAKHLW VEDHADWQQQIW EGQGINLTGAFNR GPALNALEIDSSY QGWGLLFPPLG VQIQKPPSNIKNSR TSVTSLPFVDTKGR SKKVLLKSSLLYQ KAGERQFSPWI TVAAEEENQAHNC SEIPFHSSIKTQOFK QKVTLTAAEALKL QSEILGYPELWFLC APEKFSKTSPOED DHIAYRYEVLETIG KCLOHKNNELVAL HQQALMELKLAL HQQALMELKLAL LGSELGYPELWFLC APEKFSKTSPOED LGREYPELWFLC APEKFSTSPOED LGREYPELWFLC APEKFSTSPOED LGREYPELWFLC APEKFSTSPO	ICNFDPCPT IGORIPOPCPT IGORIPOPCPT IGORIPOPCPT ICON ICON ICON ICON ICON ICON ICON ICON

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
15985	46353	IA	16076	28	1327	CVKMBUITI BDGGGBUVBUAV
15985	46353	I ^A	16076	28	1327	GYKMPVITLPDGSQRHYDHAV
		1		1		SPMDVALDIGPGLAKKPLVIRQ
ļ ·		1				GAAGLMAKLVDACDLIENDAQ
				1		LSIITAKDEEGLEIIRHSCAHLLG
		1				HAIKQLWPHTKMAIGPVIDNGF
		1		1		YYDVDLDRTLTQEDVEALEKR
		1				MHELAEKNYDVIKKKVSWHEA
		1				RETFANRGESYTV\SIVEENSAD
		1				DDKPGLYFHEEYVDMCRGPHV
				ŀ		PNMRFCHHFKLMKTAGAYWR
		1				GDSNNKMLQRIYGTAWPDKKA
		1		1		LNAYLQRLEEAAKRDHRKIGK
						QLDLYHMQEEAPGMVFWHND
						GWTIFRELEVFVRSKLKEYQYQ
		1		l		EVKGPFMMDRVLWEKTGHWD
		1				NYKDAMFTTSSENREYCIKPM
		1		l		NCPGHVQIFNQGLKSYRDLPLR
		1				MAEFGSCHRNEPSGSLHGLMR
		1		Į.		VRGFTQDDAHIFCTEEQIRDEV
		ı		1		NGCIRLVYDMYSTF\GFEKIVV
15986	46354	Α	16077	1692	1776	
15987	46355	C	16078	I	1341	
15988	46356	В	16079	83	373	
15989	46357	Α	16080	2	64	
15990	46358	Α	16081	1695	2293	EMISNPFIPSSILPLAKIRNIGIMA
		1			İ	HIDAGKTTTTERILYYSGYTRSL
		1				GDVDDGDTVTDFMAQERERGI
		1				TIQSAAATFEWKGYRVNLIDTP
		1				GHVDFTLEVERCLRVLDGAVA
	,	1				VFDASAGVEAQTLTVWRQADK
	•	1		1		HNIPRI\VFLNKMDKTGASFKYE
	ľ	1				VITISSLOKSGITKLCFFCVLKLV
		1		1		SQERCQASKCNKIEGQRFNQ
15991	46359	A	16082	1	2901	
15992	46360	Α	16083	1196	1740	FWLKAMPTIVTASPRWFSC/LR
		1				EIRSAASAIASTAIQQAIARPSTV
				1		IATTLLSNSHGNSSDSTSVEMK
		1	1	1		AIINKLRILRGRSRPORTRGPTPS
	1	1				SSANGTISQVNMVLKNGSPIEIL
		1				PSPSCLCTNGSSVPSNTTSIAAT
		1				SKTLFANSKDSRDHSALFTRER
		1				TLCPRAANSSSEPANHQHHKGE
						D
15993	46361	В	16084	49	398	
15994	46362	Α	16085	378	566	
15995	46363	Α	16086	1	2184	
15996	46364	В	16087	311	4963	
15997	46365	Α	16088	12	335	
15998	46366	Α	16089	1034	1400	
	1.3000	1	1	1		

SEO ID	ICEO ID NO.	Mar	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1,0.	sequence	""	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15999	46367	A	16090	1767	2973	SGARRLAG*KPGNAVSAAPFYF
13377	40307	ľ`	10050	1707	2,7,3	YARPV/SMISPMLIPC*CWKC/D
		l				PAAKROFRERTAVRCVAOFVD
		1				VGKLIPFWGLCVGIYATNSAET
		l				
l		l				FLHRVVQLLNDSVEHTLQDAIT
						IRDEDFTATPSPADEVAYIQLSG
		1				GTTGTPKLIPRTHNDYYYSVRR
						SVEICQFTQQTRYLCAIPAAHN
		1		i .		YAMSSPGSLGVFLAGGTVVLA
1		l		i		ADPSATLCFPLIEKHQVNVTAL
1						VPPAVSLWLQALIEGESRAQLA
l .						SLKLLQVGGARLSATLAARIPA
		1		1		EIGCQLQQVFGMAEGLVNYTR
						LDDSAEKIIHTQGYPMCPDDEV
		l		i		WVADAEGNPLPQGEVGRLMTR
1		l		1		GPYTFRGYYKRSSVLGKGSDCS
ļ						FGVFLPGLLPSASSVRRDSNVSI
				1		RNVVSLINSSRWKSGFTWHDN
						VLQLDGSFPLT
16000	46368	Α	16091	845	1920	YPMMRK/RLMKLWREAKALHS
ŀ					\	DPFEPWASIIEDADKAKSSKQA
		1				RGRGGFVRSSWQEVNELIAASN
		1				VYTIKNYGPDRVAGFSPIPAMS
		l				MVSYASGARYLSLIGGTCLSFY
i						DWYCDLPPASPQTWGEQTDVP
1						ESADWYNSSYIIAWGSNVPQTR
		1		ŀ		TPDAHFFTEVRYKGTKTVAVTP
						DYAEIAKLCDLWLAPKQGTDA
1						AMALAMGHPPERESGRYRLIPV
1	l	1				PASGYPPDPAPEWDQFPHVHEL
ł		l		l .		CDTANGCPLPELASPAGYTSSIN
						RVSTCVSTLAKSSYRSRVRVKR
1						LRRRFNGGVHFRRVVAVVIDQ
1	i	1		1		HCRTAFTIHWLQREFTEEIKTAT
	i					SPLEAFKRPONSLVVNALFRSN
						RNAAAAFSAL
16001	46369	A	16092	1	209	VSFMIIAALMMMGIDTGWLLV
	1	1				AAGFVGAIVIAS/ARPGRNSKRL
		1				IALISAIGMSIFLQNYVSLTEGSR
1		1				DVA
16002	46370	A	16093	194	657	CLDHILRAVOSLRLPLSVPAAM
10002	1.03,0	ľ.	10075	1	100,	RPGAAIGYTMVYGIIGMIN/FAH
				1		GEVYMIGSYVSFMIIAALMMM
				1		GIDTGWLLVAAGFVGAIVIA/S/
						AYGWSIER/VAYRPVRNSKRLI
1		1	1	1		
	l			1		ALISAIGMSIFLQNYVSLTEGSR
16002	46271	-	16004	1,	2202	GLATEMFLEKGDNGTEDGRAR
16003	46371	A	16094	3	3303	
16004	46372	A		144		
16005	46373	В	16096	1144	326	L

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide inscrtion)
16006	46374	Α	16097	282	599	IPVSISGCHLAGAGTQTASGVP
						VHPHGHLAHQRQSFIGNTKTEC
			l	l .		WPISSVPPHHW*A**TDIGAGSG
				l		DDQRVLDVWWNWRTKE*P*W
						W*PTNGLCRKVANRQAAR
16007	46375	В	16098	1	276	
16008	46376	Α	16099	812	1084	
16009	46377	c	16100	1	1608	
16010	46378	Α	16101	173	504	SADSKPVLPRTSASLVSASCSTR
				1		CGLRQASMESNNLTMVFGFSIG
	1	1				KVWMRS*NSATSLAKSRISFGT
	1	1				RRRWHSCSTN/SNTRRAMTMM
		l		l		TRPEFWRRQPCGLSIAAIESVIK
						v
16011	46379	Α	16102	ì	816	
16012	46380	Α	16103	742	1182	PAFIECSKAQENNHKGNCYQSG
	1					SLVSR*AFLIRHTRPVITNARR*F
		ļ				*RKVFHQRHGIAGTGSWRGFPL
		1			1	NLH*RQTVVAFQSRRAIHPM*S
						GKGREWHHFAVGAADIKFIDIF
		İ				WQHTEFGIRLNTRRAMTMMTR
		ĺ				PEFWRRQPFTEIVA
16013	4638I	Α	16104	550	884	KRYGMAASFHSGEITTHRTPAP
						WRMVVMSPARSMAHDLAQTA
	1					WRG/APRPLPDTLATMTPQAY/
						NSIQYDAEKSLWHNVE/NRQLD
						AQFFHMGMGFRRRTLWAGITN
	l					AQVCGWTA

SEQ ID SEQ ID NO: Met SEQ ID NO: Nucleotide N	VAASSV SPQVDV ELWPRH QRLRVP HTFPM
	VAASSV SPQVDV ELWPRH QRLRVP HTFPM
16014 46382 A 16105 2363 2856 MLTLWSSPTTTRWRV ANTLA*MTLSMAAMLS RFATLCHDLGKGLTPPI	SPQVDV ELWPRH QRLRVP HTFPM
ANTLA*MTLSMAAMLS RFATLCHDLGKGLTPPF	SPQVDV ELWPRH QRLRVP HTFPM
RFATLCHDLGKGLTPPF	ELWPRH QRLRVP HTFPM
1 1 1 1 1 1 1 1	QRLRVP HTFPM
HGHGPAGVKLVEOLCO	НТРРМ
NEIRDLARLVAEFHDLI	KDODV
LNPKTIVKLFDSIDAWR	NA VICE
EQLALTSEADVRGRTG	FESADY
PQGRWLREAWEVAQS*	VPTKAV
VEAGFKGVEIREELTRE	RIAAA
SDTIPQLNEHDIHRRRH	NKAIN
GERDQRNALNQLQEGF	NRYQR
HHKSGDQPNGKHRDIA	ACQEA
PALVEIERRCAHHHRHI	RQQERK
SEQHAARFGIANGDLV	RVWNK
RGQILTGAVVTDGIKKO	3VVCV
HEGAWPDLENGLCKNO	JSANVL
TADIPSSQLANACAGNS	SALVYI
EKYTGNAPKLTAFDQP	AVQAN
TRPLRLARNQHRAFRC	RITFHR
FTRIMWVEEIGQRFGN	HPFGRV
RIQDSLCVCGALRYISA	TFNVW
RKRAVTCVRSFGSDFTG	JSPPLR
APKGESIRRLLALREAG	IIHILAL
GEDYKMEINESRTVLK'	TEDNSY
SFDVFIDARGQRPLKVK	DIPFPG
LREQLQKTGDEIPDVGI	EDYTLQ
QPEDIRGRVAFGALPW	LMHDQ
PFVQGLTACAEIAAACC	GPFDKD
RIEGGIVVRMAKEGETI	LVLLDG
TEAKLNADTLVIADHN	KALAM
GGIFGGEHSGVNDETQ	NVLLEC
AFFSPLSITGRARRHGL	HTDAS
16015 46383 A 16106 390 636 KFLSFRPDRVCVACAER	ARAGL
CLRQCITGHPEACLAIP	VLVCVE
CWGGTDADQPVAVFPS	RFSQV
SPDRGT*QWLCWKFL	
16016 46384 B 16107 1 520	
16017 46385 A 16108 142 243	
16018 46386 B 16109 275 919	
16019 46387 A 16110 1 693	
16020 46388 A 16111 522 714	



SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16021	46389	A	16112	1	900	STCSSRSHLHLIQASPILPNYVA
						TAVAARIIADVQGISDTCVIAQP
						SPPHPGALRTLVFTTDEPETPOP
						PGAAGFOLVOKKOVOLHNVLY
1						VMTPHAFAOGTVTIYLPGEOOT
						LSVAPLKNVVQLVTQQHLRDR
l		i				LWWPGAFLTDFAAKVKALKD/
1						YPNHVMAORASGEAEADDDV
		1				AATIKSVRQQLLNLNITGRLPV
		1				KLDPDFVRVDENSNPPLVGDYT
						LYTVQRPGTEQGSFKVKRARA
						RVROTRAPLFGPKSPAPEHMES
		1				ELEFELIGPSKSTSGNPRGLRSE
						RPVLFCLYPEMPTAAAR
16022	46390	Α	16113	150	649	NALLGTCPGTARFPGTGSAPTL
		ļ				HPDASRSGRGRAACPRSAQWS
		1				PRPWQSVAGLAWLGAHWGSR
1						STSRTAETTASKRGHPRSPAAQ
į.						LPPGPRPPRAPQLQRALQSVHP
i						LSPRPAPLA/EPKLTAACWAEA
	}					VKITTVRYSDSCSRLLTDLGKP
						VQSMLLKDIRKVYNWQ
16023	46391	Α	16114	153	740	THQLFLPPHHQEISQTTWLSAA
						WPRHTCAHSLSPLQPPVPFSRP
1						HVPLPGPSKTWSDGPRLQSLPG/
		1				RPSERTPRR/CRSASGQPPHLGG
		1				EERLCLAAPSGRICHCEGDDES
		l			1	PLITPCHCTGSLHFVHQACLQQ
					1	WIKSSDTRCCELCKYEFIMETK
						LKPLRKWEKLQMTSSERRKIM
		<u> </u>				CFSDIPRHCRPMCGLVLVCAH
16024	46392	Α	16115	1	1194	
16025	46393	Α	16116	323	642	TGYARQKPVPALGWWTNVSAS
		l		1	1	TAPPIILPSITGSRLAVIKWLTVS
						AGKLSALLCASCQKGG/VGEM
						ANGNGERNHKQIGDTVLTPPQ
l		l				HYRKIYQPIAARSCIRCVCGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16026	46394	Α	16117	ī	1782	MAGIRSTLIKEKNMITNYEATV
				ŀ		VTTDDIVHEVNLEGKRIGYVIK
						TENKETPFTVVDIDGPSGNVKT
1					LDEGLSLLLSLVSIVVIMKLFYO	
						YEQRFGMRSTSLLNVIGSNTIAI
		1				YTTHRILVEIFSLTLLAQMNAA
		1				RWSPOVELTLLLVYPFVSLFICT
						VAGLLNLTTVMPEQYRYTLPV
				1		KAGEORLLGELTGAPCATLVA
				i		EIAERHAGPVVLIAPDMONALR
		ĺ		l		LHDEISKFTDOMVMNLADWET
			ĺ			LPYDSF\SLHQDNYLLVAFQPFY
						QLTDDAAVGVLIVPVNTLMQR
					1	VCPHSFLHGHALVMKKGORLS
				ŀ		RDALRTQLDSAGYRHVDQVME
						HGEYATRGALLDLFPMGSELPY
						RLAIGVNVAQAEVLNLESGAK
						QVLQETFGYQQFRPGQEEIIDT
						VLSGRDCLVVMPTGGGKSLCY
				l		OIPALLLNGLTVVVSPLISLMKD
						OVDOLOANGVAAACLNSTOTR
						EQQLEVMTRPRIGDGTGALRK
						RVAQLAFKNQPMMVGNWAQS
						ILHGGVIASALDVAAGLVCVGS
						TLTRHETISEDELRQRYRGWGP
						LIFALIICAQAGASVLLLLVACC
						VQAIKSPSPALNYTMKNSFILPV
16027	46395	Α	16118	380	1431	TALLIRDMPPYVEAYPNENLTT
						VMPEQYRYTLPVKAGEQRLLG
						ELTGAACATLVAEIAERHAGPV
				ĺ		VLIAPDMQNALRLHDEISQFTD
						OMVMNLADWET\LPYDSFSPH
						QDIISSRLSTLYQLPT\IQGGVLI
						VPVNTLMQRVCPHSFLHGHAL
						VMKKGORLSRDALRTOLDRAR
				l		YRHVDQPAHAGGKVLEMIVVG
						IATWLGFLLFGLNYSLLLAVLV
						GFSVLIPYIGAFVVTIPVVGVAL
						FOFGAGTEFWSCFAVYLIIQAL
						DGNLLVPVLFSEAVNLHPLHHR
						EQIGGNKMADRQCRKIVCATV
						RQLPEGGRGEMANGNGERNHK
						OIGDTVLKATGSKOGDGONHG
16020	46206	-	16110	1	940	QIGDT VLKATGSKQGDGQNHG
16028	46396	Α	16119	1	940	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
16029	46397	A	16120	114	1380	NRVHCRRCSGGDTGVSHLKGI
10029	40377	^	10120	1114	1500	NVKYRYELTDSVGVMASLGFA
						ASKKSSTVMTGEDTFHYESLRG
						RYVSVMAGPVLOISKOVSAYA
		1				MAGVAHSRWSGSTMDYRKTEI
		1				TPGYMKETTTARDESAMRHTS
						VAWSAGIOINPAASVVVDIAYE
						GSGSGDWRTDGFIVGISGVLKD
						GTGKPVQNCTIQLKARRNSTTV
				1		VVNTVGSENPDEAGRYSMDVE
		ŀ				
		l				YGQYSVILQVDGFPPSHAGTIT
						VYEDSQPGTLNDFLCAMTEDD
						ARPEVLRRLELM\VEEVARNAS
		ŀ				VVAQSTADAKKSAGDASASAA
		l				QVAALVTDATDSARAASTSAG
		-				QAASSAQEASSGAEAASAKAT
				1		EAEKSAAAAESSKNAAATSAG
						AAKTSETNAAASQQSAATSAST
						AATKASEAATSARDAVASKEA
						AKSSETNASSSADYAAS
16030	46398	Α	16121	165	299	GILPISEPPSNRIFACWGKPAWT
						ACCNSLRARR*RAISCCPSHW
16031	46399	Α	16122	1	388	MGSKSWVTQKGPAKAQSSQV
						GRVKQATATDTGRTRDSHMER
						PKCSKGYLCLPYMALQQHHLL
				ļ		SDVTVRGFVAGATNILFRQQKH
		l		i		LSDAIVEVRLCGSDLVLYGTST
						PKNLIWVMVHVVGHRPDRR*C
		l				CCNA/SVRQTQISLRAFGSFHM
				1		AVSGSSSVSGCSLFHSAHLTAL
			ļ	1		CFSWPLLCYPTLGPH
16032	46400	A	16123	130	414	
16033	46401	Α	16124	1	387	
16034	46402	С	16125	1	1611	
16035	46403	A	16126	318	401	
16036	46404	A	16127	519	658	YPGSQLLASKGTKLDGE*V*RS
				l		RLQKVGNNKLLQAKGACSNPT
		1		l		QES

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i i		sequence		
16037	46405	A	16128	<u> </u>	1837	FRPRQLVAAKPSVFADVQSMES
10037	40403	^	10120	1'	1637	VGTGGGAEAEGGVRSRGLCPT
		1				RMLPGLAAAAAHRCSWSSLCR
		1				
		1				LRLRCRAAACNPSDRQEWQNL
		1				VTFGSF\SNVVPCTHPYIGTLSQ
						EKLYSTNVHKEGQGSHTLREE
		1				KVPSFVTAEGVGAKLISPLKQE
		1				PLQVRVKAVLKKREYGSKYTQ
		1				NNFITGVRAINEFCLKSSDLEQL
						RKIRRRSPHQDTESFTVYLRSD
		1				VDAKSLEVWGSPEALAREKKL
		1				RKDAEIEYRERLFRNQKILREY
1		1				RDFLGNTKPRSRTASVFFKGPG
		1				KVVMVAICINGLNCFFKFLAWI
	İ	1				YTGSASMFSEAIHSLSDTCNQG
		1				LLALGISKSVQTPDPSHPYGFSN
		1				MRYISSLISGVGIFMMGAGLSW
		1				YHGVMGLLHPQPIESLLWAYCI
1		1				LAGSLVSEGATLLVAVNELRRN
		1				ARAKGMSFYKYVMESRDPSTN
1		1				VILLEDTAAVLGVIIAATCMGL
		1				TSITGNPLYDSLGSLGVGTLLG
		1				MVSAFLIYTNTEALLGRSIQPEQ
		1				VQRLTELLENDPSVRAIHDVKA
	İ	1				TDLGLGKVRFKAEVDFDGRVV
1		1				TRSYLEKQDFDQMLQEIQEVKT
		1		i		PEELETFMLKHGENIIDTLGAEV
						DRLEKELKKRNPEVRHVDLEIL
16038	46406	В	16129	1	1065	
16039	46407	Α	16130	1248	1658	PLALPEFISKSWKERLTDTTVLF
i						LKARRQSLCLQIGGDPLES/ILQ
t						AFCVLSVAKESQLYTYLHAIMR
1						TLKSSIARGARTFQSTSDGTVKS
1						RPSKIIFLPRLHDRFRFSALRHL
						DPKKLDLGDGSRSAHRPDRRFF
16040	46408	В	16131	969	1026	
16041	46409	A	16132	339	432	RYMVLSEFRSMQTGITKWKERI
		L				L*NCRKRE
16042	46410	Α	16133	3	443	HTWNSDRTMYLQRRHIFFKNC
		1		1		FWLQAGYSLGAPSWRSLVTVW
		1				QIIFSDFYRKTLCNSNSPLF*SLC
		1				QRQSTLLGTLLIFCSYASSPEAI
						NSAAPRVRSPRTDPRKARSPRG
						VGGAEGNGSRDTEPKGGRPLA
		L				TPSQGERLMAGTVTK
16043	46411	В	16134	34	1426	
16044	46412	С	16135	62	217	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16045	46413	Α	16136	1	609	MIPVSEKSKIGQLNLMLKGKKT
		i				KGKKVALVPAAVKKQEAKKV
						VNLLFEKRPKNVGIG/QDIQPKR
		l				DLH\RFVKWPRYIRLQQQTAIL
		İ				D/QFIQALDQQTATQLLKLAHK
						YRPETKQKNKQRLLA\RAEKKA
						ASKGDIPNKRPPVLQAGVKTVT
						NLVENKKAQLVVIAHDVDPVE
						LVVFLPVLHCKMGVPYCIIKGK
						ARLGHLVHRKT
16046	46414	Α	16137	1	824	GTRPKMPKGKKAKGKKVAPAP
						AVVKKQEAKKVVNPLFEKR\PK
1		l			1	NFGIGQDIQPKRDLTRFVKWPR
						YIRLQRQRAILYKRLKVPPAINQ
		l				FTQAL\DRQTATQLLKLAHKYR
		1				PETKQEKKQRL\LARAE\KKAA
		1				WPKGTFPTKRPPV\LRAGVNTV
		i				TTLVENKKAQLVVIAHDVDPIE
		1				LVVFLPAL\CRKMGVPYCIIKGK
1		1				ASLGRIVHRKTCTTVAFTQVNS
ł		1				EDKG\ALAKLVEAIRSNYNARY
1		1				DEIRRHWGGNVLGPKSVARIA
		<u> </u>				KLEKAKAKDLATKLG
16047	46415	A	16138	1	799	
16048	46416	Α	16139	1	344	MHQTTAKTPHTSYPNNYEEHN
		1				RRANRSKNYDRIRHTTIILKRTN
1			i	1	,	PAQTITNLLKSRFNRYNYVQPR
					İ	SCRSIFIHIPGERDLNLCIAPNPP
		1				AIFPTSIGYLGPDLNNLPTYAPFI
						TTNLTGPAPTTTHPVGNLGTTG
						NPEPGGCGILIVWCCVNWACL
l						WDSVVAAPREPYMPAGLLRHR
		l				RHYRHRREDLIAVPAARQTDE
		ı				NRRPALCHEYAVGHRKIKTNV
		l				EHTEPYILPLITHSQPLNRGAFF
		1				CPGVRFLVVSVLDFEKRLAFAS
						*SCGVQLRIRGKVQGVGFRPFV
		1	l	I		WQLAQQLNLHGDVCNDGDGV
	ļ	_				EVRLAYRVLVGQPQHYPRGRP
16049	46417	В	16140	1	1323	
16050	46418	A	16141	I	1356	
1605 I	46419	A	16142	I	1524	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16052	46420	A	16143	173	2310	QQWASRTRRWRSLITTLCVDA
						RSQHDYIALSRLFHTVMLFDVP
						VMTRLMESEARRFIALVDEFYE
						RHVKLVVSAEVPLYEIYQGDRL
						KFEFQRCLSRLQEMQSEEYLKR
						EHLAAKPETVKRDWYVVDAT
						GKTLGRLATELARRLRGKHKA
	İ					EYTPHVDTGDYIIVLNADKVAV
			ŀ	ľ		TGNKRTDKVYYHHTGHIGGIK
						QATFEEMIARRPERVIDNRRGS
				İ		LRQAGYAIHDGGNNSMAKNTS
						CGVQLRIRGKVQGVGFRPFVW
				l		QLAQQLNLHGDVCNDGDGVE
İ				ł		VRLREDPETFLVQLYQHCPPLA
		l		l		RIDSVEREPFIWSQLPTEFTIRQS
				1		TGGTMNTQIVPDAATCPACLAE
		i		l		MNTPGERRYRYPFINCTHCGPR
						FTIIRAMPYDRPFTVMAAFPLCP
			i			ACDKEYRDPLDRRFHAQPVAC
						PECGPHLEWVSHGEHAEQEAA
				l		LQAAIAQLKMGKIVAIKGIGGF
			1			HLACDARNSNAVATLRARKHR
						PAKPLAVMLPVADGLPDAARQ
						LLTTPA\AP\IVLVDKKYVPELC
1			l			DDIAPDLNEVGVMLPANPLQH
						LLLQELQCPLVMTSGNLSGKPP
					1	AISNEQALADLQGIADGFLIHN
1						RDIVQRMDDSVVRESGEMLRR
1		l				SRGYVPDALALPPGFKNVPPVL
1			1	l		CLGADLKNTFCLVRGEQAVLS
1		1		l		QHLGELSDDGVQMQWGQRLC
						RMGTQVLPPTFNIEKKVQSARY
16053	46421	С	16144	1	184	
16054	46422	A	16145	85	416	
16055	46423	A	16146	96	570	L

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown.
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16056	46424	Α	16147	2	1690	CCCCPCCCPCWGRVSGSVAGG
				ł		PRQGLGLPLSLCISPGSLNKDPS
		1				YSLQVQRQVPVP/EGLCVIVSC
						NLSYPRDGWDESTAAYGYWFK
		1				GWTSPKTGAPVATNNQSREVE
		1				MSTRDRFQLTGDPGKGSCSLVI
	ŀ	1			ì	RDAQREDEAWYFFRVERGSPL
		ļ.			1	TQKPDVYIPETLEPGQPVTVICV
		1				FNWAFKKCPAPSFSWTGAALSP
		1				RRTRPSTSHFSVLSFTPSPQDHD
		1				TDLTCHVDFSRKGVSAQRTVR
	ļ.	į.				LRVASLELQGNVIYLEVQKGQF
		1			i	LRLLCAADSQPPATLSWVLQD
		1				RVLSSSHPWGPRTLGLELRGVR
		ŀ				AGDSGRYTCRAENRLGSQQQA
		İ				LDLSVQYPPENLRVMVSQANR
		l	1			TVLENLGNGTSLPVLEGQSLRL
		ŀ			i	VCVTHSSPPARLSWTRWGQTV
		ļ			GPSQPSDPGVLELPPIQMEHEGE	
	l	ı				FTCHAQHPLGSQHVSLSLSVH/S
						PSTAAGPLLLLGG*GSALQLLL/
						SRPARPPLCAGGLGRSCWRGTA
		ŀ				VRAPSRSPPAQPGPGPTAP*AS
						MEGSAPASGSAVRPGTSTGPRV
		į.				ALSSSCYQGSWSMGEDLAWGL
1000	15105	<u> </u>	14440		1570	PWELASLPCSLSVPALSSSG
16057	46425	Α	16148	3	1578	VTQLLCTSNGRRALEFALFPGR RFRPVATNHQSREVEMSTRGRF
						OLTGDPAKGNCSLVIRDAOMO
		1				DESQYFFRVERGSYVRYNFMN
	i	1			İ	DGFFLKVTGMEWGGNPCLSH
		1	ļ			WGGTLGTAYGLSREGSQGPLQ
		1	1			HKN/HPTTVSVPALTQKPDVYIP
						ETLEPGQPVTVICVFNWAFEEC
		1			i	PPPSFSWTGAALSSQGTKPTTS
		1				HFSVLSFTPRPQDHNTDLTCHV
						DFSRKGVSVQRTVRLRVAYAP
					i	RDLVISISRDNTPALEPQPQGNV
						PYLASPK\GQFLRLLCAADSQPP
		1				ATLSWVLONRVLSSSHPWGPRP
						LGLELPGVKAGDSGRYTCRAE
	ĺ	1				NRLGSQQRALDLSVQNPPENLR
		1				VMVSQANRTVLENLGNGTSLP
			1			VLEGQSLCLVCVTHRQPPSQAE
						LDPEGTGSEPLPALRPRGSWSW
		1				LVGSSGARRRVHLPRSAPTGAP
		1	l	1		RRTLLPSPCTTPRSCWAPPAPGR
			l			LRVCTAAAPPRPARPPLCAGGL
				1		GRSCWRGTAAQGLLRGSPPAQ
				I		PGPWAQTASPEPSMEGLNSGPQ
	L		L	L	L	LOL HAVIASI ELSIMEGENSOFQ

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16058	46426	Α	16149	90	1029	MLLPLLLSSLLGGSQAMDGRF
						WIRVQESVMVPEGLCISVPCSFS
		l				YPRODWTGSTPAYGYWFKAVT
			ŀ			ETTKGAS\VAPNPP\SREVEMST
			ļ			RGRFQLTGDPAKGNCSLVIRDV
		1	Ì			OMODESOYFFRVERGSYVRYN
						FMNDRVL\LKVTALTQKPDVYI
		1		1		PETLEPGOPVTVICVFNWAFEE
		1		1		CPPPSFSWTGAALSSQGTKPTTS
		1		1		HFSVLSFTPRPQDHNTDLTCHV
		1				DFSRKGVSAQRTVRLRVAY\PP
	İ			1		ETFGISIFTVTNTPDPPENLRVM
						VSQANRTVLENLGNGTSLPVLE
						GOSLCLVCVTHSSPPARLSWTQ
						RGQVLSP
16059	46427	Α	16150	270	1001	CCCTVYRAFIRKRLKRSSDSVY
		1			1	RRKNPSASYDRDAEASLAAVK
		ŀ				SGEVDLHQLASTWAKAYAETT
		1		ŀ		LEHARPEEPSWDEDFADVYHD
				İ		LIHSPASETLLNLEHNYFVSISEL
		1				IGERDVELKKLRERQGIEMEKV
	l				i	MQELGKSLTDQDVNSLAAQHF
				İ		ESQQDLENKWSNELKQSTAIQK
	1			1		QEYQEWVIKLHQDLKNPNNSS
1			İ			LSEEIKVQPSQFRESVEAIGRIYE
	1					EQRKLEESFTIHL*AQLKTMHN
16060	46428	Α	16151	60	648	LAMSLSHLYRDGEGRIDDDDD
						ERENFEITDWDLQNEFNPNRQR
						HWQTKEEKGSLQRKEFFTDLV
				ŀ		GLCFCAYRARDYSAPVNFISAG
			1	İ		LKKGAAEEAELEDSDGFLRFFFI
	1			l		LFFH*LSFDFSFLQGGNFKPSQK
			1			GFAGGTKSFMDFGSWERHTKG
	1					I*FTYMNCCDRKTLFFFC*NSSG
						IINPIEAKQRKGKGAVGAYG
16061	46429	Α	16152	237	450	LAMSLSHLYRDGEGRIDDDDD
	1				/	E/REENFEITDWDLQNEFNPNR
	ŀ					QRH\WQ\TKEESTYGVWAERDS
						DDERPSFGG
16062	46430	Α	16153	12	85	VSFLSMGSGHCIRSTRGSKMVS
						WSVIAKIQEIWCEEDERKMARE
						FLAEFMSTYVMMNIHMIVEKN
				1	1	T*SPFCPWARATVSGPPVAPKW
		<u>L</u>				SPGP
16063	46431	Α	16154	49	481	
16064	46432	A	16155	20	248	
16065	46433	Α	16156	493	728	VSFLSMSSGHCIRSTRGSKMVS
1	1			1	1	WSVIAKIQEI*CEEDERKMAREF
1	1	1		1	1	LAEFMSTYVMMVSGRAARSG
		L_				WALPGPSMTPSPF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16066	46434	Α	16157	136	306	ASSSREGLTKFAKFTKFTKIHQI *ITLTSYSIKSLEKVCADLIRGTK EKNLKVKGPV
16067	46435	A	16158	563	1216	MMFWPELQPRVCWMGLSWVN HQNGRRRKNQKHH*QKVPLL IMASK*KLLEFLNRETKASANL SETVTKPPLPQKNTAQVGASSQ TRKPNKLAPHPAVPAFVKNTCP SRPRERQTHFLRPREDRMVAL KPIKVLPPVFMPFSSPQGIRSR MPTQHLYCRWVAPKALVPTCL FTHSLSRKEEKPKMDTPFLPY SRPPSGEWLLIVLWFFNRSQRS
16068	46436	Α	16159	1	143	
16069	46437	Α	16160	40	230	KQAAQPAVATLSGPLPHCGSFV LSLFAINLVAAHSLGPHCLYEL* HSPRRSTASLREGPRLHS
16070	46438	С	16161	233	319	
16071	46439	Α	16162	290	472	EFLDTFCYIPKVVDQ*D*THKQ QHNVAIIPHVSSAQMSTFC*RT MPTISDDPWHMASMRAV
16072	46440	Α	16163	3	553	
16073	46441	С	16164	270	608	
16074	46442	А	16165	2598	2943	HISPRGSAASFLKSVRPRTHQFR TQEDRLSPGIQDQPGQPSNTPSL QKL*N*LGVMGRICSPSYLS*P WREDHLSPGG*GCSEP*LHHCT PAGVTKSDRVSKNK*INKNKIN NDNQ
16075	46443	Α	16166	2	58	
16076	46444	Α	16167	218	342	
16077	46445	Α	16168	734	872	
16078	46446	A	16169	1233	1706	CLVFWLLYYGG,QLPFHLLPHS LGSGQQSSEESAATPSLAEGRR EAAFRCGPAGLARQAWRPHAR LPGPAAPLAAPSLAPLPTGRSG RWLFMLCAASC*SPSPAMSVYP QRRLGQGGAEDSEGGPVSRGC RAPGPQRPLAAPGEQTAARTQ AICLIW
16079	46447	A	16170	598	1159	CROTGVKIYDVWFFWLLYYPG LQLPHLLPHSLGSGQQSSESA ATBSLASGRREAAFRCGPAGLA: RQAWRPHARLFGPAAPLAAPS APLPTGRSGRWLPMLCAASC'S PSPAMSVYPQRRLGQGGAEDS EGGSVSRCCRAPGPGRPLAAPG EQTAARTGASVYLLLPHSPPLL SALHIHHLLPPS

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide	Nucleotide location of last	
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
16080	46448	Α	16171	834	1185	RSLFRHHRPSAAFPRTPCSDCPR
						CSAGTMEDCWLVLGSPELR*KS
						LHERPFHWLPSPSRARCCLPRSE
1						TPAGPVPPRPGPRRARRPRGTS
						RPPPASSRGRGALGSLPASEPLP
						ARAT
16081	46449	Α	16172	78	478	VVKNSLIFLVNQVLNSRDFHVF
ľ						FKTT*NCPAQAHTAALYQGLGP
						ACSQRASHWMSSSSKRSSKSLK
İ					l .	KSA*TAFSLAAKSIKSPGAASRL
		1			l	GNGKARLCQVRFPKEQVISALQ
İ	ì	1				GHYPHKKRRWVRGNRLKLEST
		ļ				F
16082	46450	С	16173	1323	1656	
16083	46451	A	16174	77	231	RSCMTLMWPRSTP*FGLMERR
	1	1				RHMFDWLLITMLWMLPTKLGS
	1	1		l		SKLSPAA*F*SSCLIL
16084	46452	Α	16175	1	255	STSKGEKPAIRAMAHFLEPHPIN
1	1					FLGARARKADLSPT*YHFFKHL
						DNFLQGKCFHNQQDAENAFQQ
						FFKSQSTNFYAGSSFVTEV
16085	46453	Α	16176	690	1020	SPVLQRAFKNGPGTANERTVQ
						WWFKKLCKGDESLEDEEHSG*
		1				PSEIYNNQLRAIFKADPLTITRE
l						EVFPHPLYSPDLSPTSYHFFQHL
						DNFLQGKCFHYQQDAEHAFQE
16086	46454	A	16177	1	383	MHRSCGSDAYALLGERSVSLR
	İ	ľ				RCAVRHAGGIRSANADISNDKA
		į.				GEKPARRKTKGSCPTLIGAGLV
						QSLSPIQSAADAGELGSNGILPD
						VESLRIPVQYLANLLTAGDTKP
		l				VLRALKRMLAMRHYKRAETV
		1				DGKVDTRALEEVGLTEAQAQE
		1				MYRYLAIANYEDRFVVPSSHRE
						LAREAFPEKNGCGFTFGDGCH
	1	l				GSDTKFNLFNSRRIDAID*PCPD
		l			1	*RWTGTLGLPASGLFTRFIVTY
	1	1		I	1	VSIRTSDTSSMPHSTPSQA\TERS
1	1	I	1	l		PNNA*ASLQTC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16087	46455	Ā	16178	740	1977	SASASRSLEHITVTPSTGILTCFP
						STTPFGLAL/GGTLGLPASGLFT
				ŀ		RFIVTKSSIFSVTPLVQPKPSCN
				l		GYSPPPPPVPHRREFQRRVIAEK
	į			1		IFCQHHRAKLNRLEERQYLTVN
		1		1		FRAIAQRVNIGITGLQITIDANA
l				1		AINROPGASCOGRFPGAVRCGG
				1		NYRIHFDIALAVORWPECONPF
				1		FPAPVTEISTADERPSPLTGCLQ
		l		l		RLVTFGSVIFGIINMIMRLAPIGR
				1		FGAMAFTIGKYGVGTLVQLGQ
				1		LIICFYITCILFVVLVLGSIAKAT
1		1		l		GFSIFKFIRYIREELLIVLGTSSSE
				1		SALRRMLDKMEKLGCRKSVVG
1				1		LVIPTGYSFNLDGTSIYLTMAA
1				1		VFIAQATNSQMDIVHPITLLIVL
				1		LLSSKGAAGVTGSGFSVLAATL
						SAGGPIAGSGSGADPRYSTALC
16088	46456	A	16179	757	1037	FLDPSPEPNVTVTPSTGILTCFPS
10000	10.00	ľ.	1	l'*'		TTPFGLALGVDSPCPD*RWTGT
				1		LGLPASGLFTRFIVTYGVTPALD
		1				TGLM*LRSCGPPRSML*HRTQQ
						KC
16089	46457	С	16180	1	313	
16090	46458	Α	16181	178	290	LLLGLASEN*HCPIASEAP*TTT
		<u> </u>				DTELWVTLTVEGKS
16091	46459	A	16182	1	251	
16092	46460	С	16183	129	1136	
16093	46461	С	16184	117	748	VII OI I I PROPERTIES I PORTE
16094	46462	Α	16185	636	748	LLLGLAAED*RCPIASEAPQTIT
	14140	١.	15101			DTELWVTLTVEGKS
16095	46463	Α	16186	37	444	DPQVGKLLQHCPARIKMTVTF
						NNHRKPQLSKFHQLYSEDTQQ
						QIIRETFHLVFKRDENVCKFLGT
				i		IWSLTTGKLNRGQAPLFGIFHT
				i		VELHLEIRSSCQGIVHF*LTCFSS CPCSGPFVNSPFYLPLESGHSK
1.000		l.	16187	179	422	
16096	46464	A	16187	1179	422	LQLGPKEKQSRVLGSFFQVPSC
				l		GKMGCGSSGSRHIGFPP\PQLSG
				I		GPLDPRQGSGISEMK*TE*GGY
15007	16165	L	16100	1064	1274	PKGSPQHYSRKHQGS
16097	46465	A	16188	1064	1374	KEQCLQALSSVKNREKRPSSFH
				ĺ	1	PICWSAVSFSVGREGDMWPLAP
		1	1	ŀ		FPDSENVAGAGALLLRLAALPG
		l	l	1		LSWALRPPPPAPVRSWATSGW*
		Ц.	L	l	L	GGQGREPGRGIPPRW

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16098	46466	A	16189	44	515	VGGGGVQPSMTMGDKKSPTRP
10070	10400	ľ`	10107		1	KRQAKPAADEGFWDCSVCTFR
						NSAEAFKCSICD\VRKGTSTRKP
						RINSQLVAQQVAQQYATPPPPK
						KEKKEKVEKAGQKRKPEKRPR
						KFSP/SVVTKKNTNKKTKPKSDI
						LKDPPS\EANSIQSANATTKTSE
						TNHTSR
16099	46467	A	16190	382	795	DSGRSWKRLLSPPRPMCDTWA
10022	70407	n	10170	502	1,75	LGSYPRLCFAVSPPFSCSLPLPSL
		l				FPHSPLLGRPKRQAKPAADEGF
						WDCSVCTFRNSAEAFKCSICDV
					i	RKGTSTRYLKICVTWVVVVI*Q
						VFIEHTAICALGVQWYKDRPGP
		ĺ	}			CSYGA
16100	46468	١.	16191	2	343	CSYGA
16101	46469	A	16192	472	625	KTSLFOHWLWKVTIVIVMF**L
10101	40409	^	10192	472	023	RVNVGKLLNNIHRYNSENLASL
						GSOWKFP
16102	46470	١.	16193	343	693	KTSLFOHWLWKVTIVIVMF**L
16102	464/0	A	16193	343	693	
		1				RVNVGKLLNNIHRYNSENLASL
			1			GSTQIPLKAFTNPCHTSFTLCKH
		l				MVIQAHQEKQLIQQILYLGDLL
						ET*HFQAFWQALDENMTLLESI
16100	46481	_	16104	240	534	TGFEGF
16103	46471	Α	16194	340	534	TAIMNDMVTI*TRKFMTNQRLQ
		ĺ				RKQMVIDVLHPGKATVPYLILI
		<u> </u>				EQLRFARHCAWYGGDSDETSH
16104	46472	A	16195	80	552	PIPHLEKFHAPTRRTFRGNQM\V
						IDVLSPPGKADQCP*AQEIPGKT
						*PKMYQDPHPDVI\FVFWIPELH
			İ			FGSRGKTKLAFGNDFMTSLGFI
						AKEKWNPKHRTLQGHGPGME/
						ERKKTSKGKPTERERQGTGLEG
						KFQGGLAKGPFFGAWAKRRNE
		L_				VSSS
16105	46473	A	16196	248	474	YPETPESPTQLSSGQPHREQPPE
				l		MPESPTQPSSGQSCQEQPPETPE
						LPM*PCLAQPSQKHMIKG*QNN
		_				KHHENKM
16106	46474	Α	16197	223	669	PRPSLPACSPWRIVDDCGSLKS
				l		WKRVCQWLQPCPNALP\$CLGS
				i		PLRCEFSGFPRN*LGLGSTTGSF
				l		AVWGGLFSTIDCGLVRRPGPSP
						FHPVLPALTSVSLPPTPPGGPLA
		1		l		MVGSAMMGGWSGEWEILHLA
				l		SSTHPYPSLPSPAPPFLE

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence.		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16107	46475	Α	16202	188	484	HFGRVEVGRSQGQEFQTRLAN
	l					KAKHHLYSFMNESQISPIGC*P
	1					M/PPKLIRKVHREKNCAGRNLT
	1			i		R\QRSGFKTGPPGPPARPSSPTLP
	Ì					PPPPLPRLPRRP
16108	46476	Α	16203	36	409	
16109	46477	Α	16204	651	893	NTSVKNNFSFMNESQISPIGC*P
						MAPKLIRKVHRERNCAGRNLT
						RAEKRLQTPGLPVPPARPSSPSS
						PPTPPLPAATPGGR
16110	46478	В	16205	13	441	
16111	46479	Α	16206	3	205	NSSNSFSIQFCSFAGEELCSFGG
					EEAFWFLEFSAFLRCFFLIFRDL	
						STFGL*CAPLEEKRHSGFWNFQ
						PFCAAFSSSSGIYLPLVFDAGDL
						QMGFLCGCPFC
16112	46480	Α	16207	23	841	EKAKEFRRAEEKKKEVPAVPET
						LKKKRRNFAELKIKRLRKKFAQ
						KMLRKARRKLIYEKAKH\YHK
	1					EYRQMYRT\EIRMA\RMARKA\
	1					GNF\YVPA\EPKLAFVIRIQRVS
	1					MGVSPKGSERCCKLLRPFVKIF
						NGNLL*KLNKGFRLTML\RIV\E
	1					PYIGMGGTPILKSVNELIYKR\G
	l l					YGKIH\RKRIA\LTDYALIARSLG
	l					KYGIVCMEELNHEIYTVGKRFK
		ĺ				EANNFLWPFKLSSP\QGG\MKK
						KTTHFVEGGDAGNREDQINRL
						HRRNELTRLTWIFF
16113	46481	Α	16208	1137	1200	PPSAPAPG*LRPPALSSGSP*PPS
	1					APAPG*HAPPPSAGTPPP

SEO ID	SEO ID NO:	Mar	SEO ID NO:	Nucleotido	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16114	46482	A	16209	1	1480	MNECRGVEITTVCSEDWIQRHL
	i					PGAVVPLKGEGNVGVATEDHY
	i .					QEDSFGFGLRGTGKPLKLEKSR
						PGAVTHTSPSDGLTQYVPDSLT
						AMPAAAVVIWVIRPGCGVOOO
						IGFLRHSNIDDIRPRTIIRAVAPV
						KLHGIRIKMASFTSATRLTDEO
					l	VITAPADSGIVIPRFSDNWVAG
						GKTHLTHKDSHKLKVKGWKK
	1			i		AFHANGHQKQAGVAILTSDKT
	1					NFKATAVKKDKEGHYIMVKGL
						DQQENITILNIYAPNTGAPKFM
			ŀ			KOLLIDVRNEIDSNTIIAGDFNT
						PLTALDRSSIOKVNKETMDLNY
						TLEOMOLKDIYRTLHPTTAEYT
	l .			l		IYSTVHGTFSKIDYMTGHKTSL
						NKFKKSEIISSSLSDHSGIKLEID
				l		SKRNHONHANTWKLNNLFLNN
						RWVNNEIKMEIIOLFELNNNND
						TTYONLWDTAKVVLRGKFIAL
	l			l		NAYIKKTERAKKTIYSHTSGI*K
				ŀ		QEQTKPKPSRRKEIIKIRAELNEI
				•		TTKKTIOKK
16115	46483	A	16210	32	398	ASERLGGPGSPG*/PVNPRTDGS
	ļ				1	GAAWLSQRAQCTQSDPRV*EV
	ŀ					PEHLSPSTPSQEYRGLPAL\PHTS
				1		RPSHCLPACPLPSRTOPWVPVK
	1	l	ŀ			PGPTACRGFLQHSPTPGSPSP*S
	1			1		FOGSSWWDTDL
16116	46484	A	16211	2	681	AEPWDEGGDGVPCRVPSSGSV*
				l		V*TLYLDSSVQLCGTEAPQHPV
				l		DGPPGGKLCHEGSLHLRQGQIP
				l		GPRLVLQFVQQAGVCGPGGNA
		l		l		VYSTPRDSASLVSGRTSMLFLR
		l		1		KFGKTWRKGKLHORGPPASPV
	l .					LGRECNWG*VIGRHPAOSS/SGP
	1	l				ACAKDAAELGAAMPGLVSGPL
						GAALQPRGAGVQLGAATGPGS
				l		CSK/RGSPKADAR/PRHSOEPGO
	1			l		VPPHAPQGPGLS
	1		L		L	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	Į į	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		_				
16117	46485	A	16212	102	1005	VQWSARPDATPLSTEPSLADYP
						EAQSHIPRRGPRRPEKQGRGGK
		l				PPRALGRERAGGGSRGPPASPD
		ı				PLSRPCSQAPPPGASAGA*DAP
		l				ACRREAGSDGVPCRVPSSGSV*
		l				V*TLYLDSSVQLCGTEAPQHPV
		l				DGPPGGKLCHEGSLHLRQGQIP
		1				GPRLVLQFVQQAGVCGPGGTD
		l				GSGAAWLSQRAQCTQSDPRV*
İ	l	l				EVPEHLPQVDTHKGELYQ*PGG
	i	l				RDGPGWEEPPQCGMGVSEAHG
	l	l				GAGGPGEAGKGRLQRKFGKT
	Į.	l				WRKGKLHQRGPPASPVLGRPR
		1				PE*VLLRCD*EG*PPSAGQLL
16118	46486	Α	16213	378	494	GTACLQIPQTQGV*QRGFLAEY
1		ı				IQWQLRMRSGSIPVRTG
16119	46487	Α	16214	18	894	SVGSLCCLQETPGLRAFGVPWR
		l				SAQRRWPRRSAASLLSLVLPGR
						AHSPRPTLAPGPLEAQQPPPGG
1		l				SALAQHSRTAPAGTSAACLWP
ĺ				l		VSEQGAAPMCQVTRPPGSQAP
						CSQPSAHAHKHLTTGSSSPAEA
				l		SALPRA*TGSRWETKSSERVSSF
				l		SA*GTACLOIPOTLRRRTG*RO
				l		VGRRLPGSGGFDGCSIFS*S*PV
1						TYSLRSVSATSSNEATOLGTT*S
}						NSIIFEFCSAQSTAVRCRLFST*T
İ						DAPWSRSSSTTKTCFSITASCSA
		l				VLPLLLEALMEAPCPCLACSSA
1		l				s
16120	46488	Α	16215	340	582	
16121	46489	В	16216	93	711	
16122	46490	Α	16217	1378	2081	LSRKKPNWPSSMPWYQFSLIWL
		l				MSVKTSPSWKPSSPWFSGWKS
		l				YSTLOLGWSRGGMLLLGVOM
		l		l	İ	DLLEDWKGDEAAILGTVVQGK
				l		PPLDCSGSPRSK\APRTWAYTPA
1						WGVRKLPASCA*ASLPARPGSA
1		1				WPASGRSAAPPP/AARS*LICSM
l						SVSRRLCMLVSATSLESSNFSSA
		l				WRSSSSGILPCRFSMSRESFTSA
		l				RCFMDSPASSSASLKLDSLPSSF
		l				WWRIRHSLRSPSDCG
16123	46491	A	16218	191	288	
16124	46492	A	16219	67	209	
16125	46493	A	16220	96	341	
16126	46494	В	16221	1	1185	
10120	40474	D.	10221		1105	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16127	46495	Α	16222	1192	703	AMPSCTRAGLAFVCHVRRRGC
				Ì		LSCRGVIPASSOCLFPAPMGTTE
		l				ATLRMENVDVKEEWQD*DLPR
		1				PLPEETGVELLGSPVEDTSSTSN
		l				TLNFNGAHRKRKTLVAPEINISL
	i	l				DHSEGSLLFDDFLDTLDDLDIN
		l				VDDIETPDETDSLEFLGNGNEL
						EWEDDTPVATAKNMP
16128	46496	Α	16223	8084	8251	TAGDERHGPSRVRMFRIKGRRL
						GWMGPLGRGKR*VTRGMWPG
						KGPAYPGSSASSF
16129	46497	Α	16224	645	1317	WGASRLREPPSPRTLHSPRPGP
		1				ASAPPLRPLPQNRTPSPSAASTT
ŀ		1				APPQPSEASRLPA*KAKPKPHPP
ŀ						SAPRRGGSPPRHSAQPACGRRR
l		1				PRSVGGAAPYAAERRNSAPQGP
						PQRLRRTAEGGKRPSNPRQRSH
				1		PRQRGDSLP*LGPWSRDRRAGR
ľ				1		DGRDGSAGGTRRSPSPRYASPE
						PLGPAAPASPAGASWTGGPGRR
				[ADDSWRASVFGAQSSACFPRA
16170	46400	В	16225	72	378	RVL
16130	46498 46499	A	16225	72 151	372	GGQHWAKAWRQTLGPEMPER
10131	40499	l ^A	10220	131	372	TGACGCGSL*PFTTPPPPAICVD
						GTFHKYVFTPDGNCNREAFDV
						YLDICDDDDF
16132	46500	A	16227	182	1083	HPGPCPDHEOVGSMGLVEDPV
10152	140300	ľ.	1022	102	1003	SPGLPLTLNLGPDNPPTCPAVLI
l		1				WDDAREGKD*EG*SSLGIPPPP
		1				HGHPVLCDTHRIVIVLKNROLC
		1				PF*DAASSSAAHPTPHLHTLGL
		1		l		CELPCGLLAPSLHTTPTSDSSLL
						PODLASTKPGTSSAPFTINAHOS
						DIACVSLNQPGTVVASASQKGT
						LIRRVCMVGW*PHTT*HVGITA
						DLSSSINFSHDSSFLFPTLPCPCP
		l				PAHLPYLGLASAACRLARVGK
		l				VGPMIGQYVDSQWSLASFTEM
		1				PERTGACGCGSL*PFTTPPPPAI
	1			1		CVDGTFHKYVF
16133	46501	Α	16228	202	517	GVHCHGFSAGQAAGQEPTSVH
					1	l
i .		1			l	ETLLPHQLQDQSALRGGVQNR
						ETLLPHQLQDQSALRGGVQNR QRHQAAEGPGERAGAAVSVGL

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16134	46502	A	16229	141	633	WENARPEPSQGDLRAASVVSPR
1				İ		LPPRKGQGGLGYFGRGAGHAH
						GTAGQGASESETLLPHQLQDQC
				i		ALRGGVQNRQRHQAAEGPGER
				1		AGAAVSVGLGEPQCH*VGAGR
İ						AARGPPILEVPAGGGDAAAECP
İ		l				AGPHGQNARGQQSGISQLRHE
						REFLLRPFPQAPQL
16135	46503	A	16230	65	466	VVFAEAAEASLMTSGVGGGAA
						QSPRSRRGGWWPLGEMPYANQ
	1	1				PTVRITELTDENVKFIIENTDLA*
					į.	GYRGKRLAALGRGRPKPGPRG
						RGRGCSGWGGGDVLPELPPCT
		l		1		RALALGACGAGPACRGCSAFTP
1		1				LGF
16136	46504	A	16231	42	491	SGGKEAQGQSGTDQTVVVLSN
		1				PTYYMSNDIPYTFHQDNNFLYL
	1	l	Ì		1	CGFQEPDSILVLQSLPGKQLPSH
		l				KAILFVPRRDPSRELWDGPRSG
		l		i		TDGAIALTGVDEAYTLEEFOHL
		1				LPKMKGNKWEQKSHYKPDWD
1		1				*NLSCLFRQVLNFVIVAPP
16137	46505	Α	16232	531	976	NOSIVRVNVDNSLSIPVIAAELG
		1				VRKPSEKGMQQKKKTKDLGFR
		1			i	AGKESKTEWRK*GLODMASO
						MFASPLK*PVTAAFHDSSMPSS
						LLQIEMEQLFLEARLQLDSKSE
						ARRNOCDSMLLRNOOLCSTCO
						EMKMVOPRTMKIPDDPKAS
16138	46506	A	16233	2	419	SFLLSSSYTRTKSILRWPSLKKR
10150	10000	-		_		ESGFPLNSSNSGVRKSC*LSSDP
1						KSGTIVCPRFSNTISPSSSSSSSSS
		l		l		CSKTFL\PSFSCLPQKAAPNLAR
1				1		SWGLPGSRTARLSQGPDRGPRG
						VCRLGSGAAEPSORRGKPSRDE
		l				iv
16139	46507	A	16234	27	792	INSIRILSTCKMGHPRFAPQAPG
10137	10507	ľ.		Γ΄	I	PLRTSEKGPAVLPAWGNSSSDH
				i		SSEPSVGFYFNEDSDGSHPVLQ
						MRACSPDPAQGARPSAEAPGQP
			1			*HSASRGSDPKVASEWRLPHAE
			İ		1	ASPOLPGFSGAASPAPKHSCSSF
			1		1	SCLPQKAAPNLARSWGLPGSRT
1			l		1	ARLSQGPDRGPRGVCRLGSGA
			1		1	AEPSORRGKPSRGTHRPEPRAG
					1	GRLSPAERSPETEHAOCGRRGS
			1			LRND*TSPVLRGAQKEVGRRD
			1		1	CRGSSAILRAGEL
	L		L		1	CKOSSAILKAGEL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16140	46508	IA	16235	343	969	YDKTRLFNTVPLPPPHKKIIKKII
		1.				ODIKGEILRAFPLRTGTKHRCPL
	İ	1				SPLLFNILLEVLARAIRQEKEIKS
						IQIGKEEVKLSLFADDIIIYLESP
		1				KYSSRKLQELIKEFSKVSRYEIN
						VHKSVALLYTNSNQAENQIKKS
1		1				TPFTIAAKNKIKYLGIYLTKDAK
		1				DGYKENYKTLMKEIIDDKNKQ
		1				KYIP*SWKGRINIVKTTILSTAIY
16141	46509	A	16236	15	403	ICESRRGRVCSALVRRYVGEGF
		1				GKAGSICMPOAPDKSSRWTILI
						RPPRFPPS\PSCSPGKEAPGGSEL
		Į.				/SPLPHAWPSDPVSPLGDRVTA
						NPAEIR*EPGPOSAAPGFREORR
						RGQMG*IWYLTPTPPGGQ
16142	46510	A	16237	19	454	KGKSLLCTSPKVCRGRLRKSRK
101.12	10010	ļ.,		1"		PETKVTHICMPQAPDKTSRWTI
		i .		ì		LIRPPRFPPS\PSCSPGKEAPGGS
		1				EL/SPLPHAWPSDPVSPLGDRVT
		1				ANPAEIR*EPGPQSAAPGFREQR
1		1				RRGOMG*IWYLTPTPPGGORPS
i i	1			1		SAASPRGFPLV
16143	46511	A	16238	276	898	MKGTAPWRCGSRRGRVCSALV
10145	1.0511		10230	1-1-4		PKVCRGRLRKSRKPETKVTHIC
					l .	MPQAPDKTSRWTILIRPPRFPPS
		İ				SILOSWEGGTGWLRTLPLPHA
1						WPSDPVSPLGDRVTANPAEIRIL
		i	ĺ	l		GEDPESPEETLNLILRTPIPPKPR
						GNAKSHP/GGHRSHR*EPGPQS
		1				AAPGFREORRRGOMG*IWYLT
		1				PTPPGGQRPSSAASPRGFPPTNN
						SSRLPAEPE
16144	46512	В	16239	177	284	
16145	46513	Α	16240	253	416	CCPVLPGPWAAGHPGD*PDGP
		1				G**AQPHKECRGGFAPLQVTLG
		1		l		PKQPSGKRVNK
16146	46514	Α	16241	1401	2246	NIYLLLQVFDCWGLQGILGVSN
		1			1	PSYSIRRCYTAGQSPRGLPDPGR
				1		F*G*EESTGEQADCP*GSHRRSR
		ŀ		ŀ		SAAGAADGGLDPGAHTITCGC
l		1		l		LSHRGRLVPAQKSSAAL*APSG
		1		Ì		AKPAPTVAPVHDPARQG*GGQ
	1	1		1	1	ARGQGARQPHSHREGLCEC*A
1	1			1	1	VTTL*EPHLRNSSAPA*ERGPQG
	1			1	1	IRVPEGQSGDQESSAEEHHGGD
1	1			1	1	PGGGKPRCGQHQEPLGAGWPS
	1	1		1	i	PAGVEPLLGGLQKGSDGAP**E
	1	1		Į.	1	PORGCVDEAQOSSPGAVPEAK
1	1	1		1	1	AIAVOPPAPDVFAAVARCD
	J					P.III. C.III. C.IIII.

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino aeid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16147	46515	A	16242	30	461	IPAETTC/EPAKHFHCFTDGSSN
						GKASYSGSKGKVFOTPYISAOK
		ļ				VELVAVIEVLTAFDMPINVISDS
		Į.				SYMVHSTOLIENAOLRFHTDEO
						LIIKTKKGGNRDYGTAQTQLNL
						ALLTFHFSSLPKGQMLPAAEQH
		1		l		LQKPAAKTQAE
16148	46516	Α	16243	I	573	
16149	46517	Α	16244	1	1689	
16150	46518	A	16245	1	1416	
16151	46519	A	16246	27	779	KNHFNWKDKFPPRLRISFLMVA
l						TVKPPEPTPLKRLTEKPIWIEKW
1						PLSK*KLEALEKLVTEQLENGHI
						APTFSPWNSPVFIIKKKSGKWR
ļ		1				MLTDLIAINSVIQPMGALQPPGL
						PSPAIIPKNWPLIVTDLKDCFFTI
						TLAKQDCEWFTFIIPAVNNLQP
						AKHFHCFTDGSSNGKASYSGSK
	ļ					VKVFQTPYTSALTAELVAVIEV
		1				LTAFNMSINMISDSSYVVHSTQ
- 0		1				LVENAQLQFQTDDTNDFIYPVA
						NSS
16152	46520	Α	16247	2	394	
16153	46521	A	16248	63	1123	ELDVMGLPEERVRSGSGSRGQE
				1		EAGAGGRARSWSPPPEVRLVSL
1	1	1				LRLNMSQKINDIWASISVAEFW
1	1	1				GDIAKEFYWKTPCPGPFLRYNF
	1	1				DVTKGKIFIEWMKGATL*LCLF
						LVWGRSYTPNHWFFLVCLREG
		1				NEPGGQLDGREQVAARRVHGS
		1				YSVLLPAGIQKGDRVAILSNPFL
		1				HSLNPWGFLLSLRLSSLP*GPM
		1	1			DIVPTRQHGAY*GLECLLLPKD
		1				MDMGGSCQVPFEPTRLPL*AFL
		1	1			FSRGFPVRCCIVVKHLALTLSSH
		1				SLYTWTLQGQGTGVSCPHLELF
						TVLPCPSLQISWNQGIDLWWHE
		1				LMQEAGDECEPLPSLHLVVSRG
		1				GHKTLTCSPFFPQGVVHTVGGY
16154	46522	Α	16249	35	219	ALGVHFVLGPVGDGSRAVDCG
		İ		1		QQL\PTSSACTTSGLSISRRCSGI
1 _			l			WRMNQKKQVRGPSPKE

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=linknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16155	46523	A	16250	1934	3213	WPAPPSQVPSLDYGLALWFDA YVSVHOTOETSQCSPESSLTPRI LCGLRILOPY AERIPVVATAL INFTLRGYAWPNGALPASTVPC GFAACGEFLCSVNGRQAPAGS SGFLPPYBSLCPHTVCRATFQCK EDSTCISLGRPPWGSETHLLCPA PIPSVPGVPCGTFTFQCEDRSCV KSLSASPLPLLCPPHHLPSPD CGLOGPSSRIVGGAVSEGEWP WQASLQVRGRHILGRGSFLPPH LTPPHPTPPHPSMASTVLWTVF LGKVWQNSKWPGEVSFKVSRL LLHPYHEEDSHDYDVALLQLD HPVVRSAAVRPVCLPARGRIEN RGSRLLRSHFCP*PRTGPISNAL QKVDVQLIPQDLCSEVYRSSPG KGGALTLLQSFLLSPFQGDSGG KGGALTLLQSFLLSPFQGDSGG KGGALTLLQSFLLSPFQGDSGG
16156	46524	А	16251	511	853	CCWNMALIQTFQMSMEIPLCTT LS*TALHLASANGNSGVVKLLL DRECQLNVLDMKKETALTKAV QCQEDECALMLLEHGTDPNIPD EYGNTTLHYAIYNEDKLMAKA LLLYGADIESKNKVEEEMKH ESNNVGLLENLSNGVTAGNGD DGLIPQKKSRTPENQOPPDNESE EYHSHTETELFSRIYP
16157	46525	А	16252	1050	1350	CDPQMCCCFRPEAATPG*REAA SAGPWDSAPPSPARTSLPSAHSS DYPLPRSRSSSASAASRSPPRPL STIARRLNPRRSGRSLPSGQPAG PSDPRGFCL

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide inscrtion)
				- Sequence	1	
16158	46526	Α	16253	2	3486	SSQSGAAAGAFCGLCVDLELTA
						SCDLEADSEDRTEHLATLQLRP
		1				DFSGSRCPAPQKHLQTHRPHSK
		l				PWPIY*PT*RSYLAISPHSW*AP\
		ı				PPSRRRAPPAFHPHTQACPSTCY
		1	İ			CHTLASRRGPSNGRYHRPGYPH
		1				PTAVQRDPPAGPRGCQSPC*HQ
						PPACRRPCGRHYR*HGQHDPPP
		1				WQ*HC*SRSPCHPPSQRDAVTW
		l				PFIHIPAED/SHTQPQKSP\PAFRP
		1				HTKTCPFTCYCHTFAGRRGPSD
						GRHHHLGYHHPSAVQWDPPAC
		l				*RGCWRPCQREPQTCPCFCGR/P
						PPGTSLA*PPHPRVPLKEKLCQY
						DEFSEGLRHSAHLNRHQRVPTG
						EKSVKSLERGRGVRQNTHIRNH
						PRAPVGDMPYRCDVCGKGFRY
1		l				KSVLLIHQGVHTGRRPYKCEEC
						GKAFGRSSNLLVHQRVHTGEK
						PYKCSECGKGFSYSSVLQVHQR
						LHTGEKPYTCSECGKGFCAKSA
						LHKHQHIHPGEKPYSCGECGKG
1						FSCSSHLSSHQKTHTGERPYQC
1						DKCGKGFSHNSYLQAHQRVH
						MGQHLYKCNVCGKSFSYSSGL
1						LMHQRLHTGEKPYKCECGKSF
						GRSSDLHIHQRVHTGEKPYKCS
						ECGKGFRRNSDLHSHQRVHTG
						ERPYVCDVCGKGFIYSSDLLIH
		ŀ				QRVHTGEKPYKCAECGKGFSY
						SSGLLIHQRVHTGEKPYRCQEC GKGFRCTSSLHKHORVHTGKK
16160		١.	16254	222	695	
16159	46527	A	16254	333	695	ISVFRSPGQSTSQHDAATWPFL
						HISGEGPTPSRRKAPPAFHPHTQ
						ACPSTCYCHTLASRRGPCNGRY
		l				HRPVYPHPTAMQRDPPAGPRG CQSPCWHYTPACRHPCGRHYR
		l				*HGOHDPPPWQ*HC*FGSPGQS
		1				
		1				TSQHDAVTWPFLHIPGERPTAS
1				i		RRKAPPAFHPHTQACPSACYCH TLASRRGPSNGRYHRPGYPHPT
1		1		1		A VQRDPPAGPRGCQSPC*HEPP
1		1		1		ACRHPCGRHYR*HGOHDPPPW
	l	1				0
16160	46528	A	16255	9	437	SCNMSLCRCNVCIRAAFKEGTP
10100	100220	l^	10233	2	,,,,	ALLVYSAILSFTPVSLAKSCSISI
1		l		1		AIVSNILSWLDPFKRGVVSELSR
		1	1	1		LPSRVLL*L*LVHSLLKETSFRS
	l					SSFSSCRACFLRSSQ/STAGPCR
				1		HLCRPTSAGIGEORRCRPESHH
		1		1		EIVDRGI
	<u> </u>	1				DIVDROI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16161	46529	ĪA	16256	619	1138	SCNMSLCRCNVCIRAAFKEGTP
		1				ALLVYSAILSFTPVS*DPEAFSP
		1				HLPSNCWSLPPSLSSNVSRDRG
	1	1				AEAMQAGKSPWCVFPP*EPGPT
		1				GCEPPRTLPPAPFKLCSL
16162	46530	A	16257	408	801	SQRRGSVCRLVHGRHGQPGKG
		1				TSPHSSLWDWQPGPQPSGPPWF
		1		1		EGGASPGTPSLIPRS*SASCCFP
		1				WHPGCSC*GAPASQC*VALSPQ
		1		1		PWFPLPCLSVPSPVGAQATGPK
		1		l		HVRTPPGSDSAPPQLQPQSRI
16163	46531	Α	16258	1	199	
16164	46532	A	16259	665	1006	IGSPDGRNVIRERTVTNETQTEG
		1				EREKEEERESFHTGDVLKVGDL
		1				GPQGSSREFQLLSNLTVPNFIDI
		1		1		QFPK*VSAAC*DGPSHTHSSYIS
						SSLFIIRASQSSPGDFSISVRY
16165	46533	A	16260	247	404	
16166	46534	В	16261	258	934	
16167	46535	Α	16262	519	924	NRTIQADLLYPDDAGNPLPARN
		1		1		TRQASASADPPGARAGERRRG
		ı		Ì		AAGPSPGR*QQAPGPPPPGEGA
		1	İ	!		GARPG*EGDAAGGATRFVGED
	1	ļ				ARLLRAGCHRGGRGEAEEPLQ
		1				ALRHPGLQGLL*EGAPHGELAL
16168	46536	A	16263	121	214	VAFAHL
16169	46537	A	16264	139	272	
16170	46538	В	16265	1	1962	
16171	46539	Α	16266	2373	2554	SATPWLPTGLAAPTPGLAAPGT
						RSPPGQT*PERSPAASPKRPERP
				ļ		ALDFOPPROPPPPP
16172	46540	Α	16267	919	1653	WNCFVCSADTMAQIMAEQEVE
				i		NLSGLSTNPEKDIFVVRENGTT
		1				RRPLQPPPRGRAS*GGAGRLCS
1		1				QLITEQADIALTRGAEVKGRCG
l		1				WRPLGLRTQTLRGRLKRTSPAQ
		1	i			QESHNMSKGPEATCQLSGTLG*
		1		1		EQ*GF*LLLWALAVAGKHTAN
		1		ŀ		SHHLSALVTPAGKSYECQAQQ
		ŀ			1	TISLASSDPQKTVFCRTYDSGCF
		ł				DQSSLLFFFPIAEHKCPVDEREQ
		L	L			LEETLPLILGLILGLVIMVTLAIY
16173	46541	Α	16268	2	321	PAWPPWPGITGFCPQQDSPSNT
		1	1	1		FVVPAAHTV*CSHPCSYYKGIR
			l	1		QMVGLPQPLFHPDCPVTPQAHT
		1		1		DSLNTLGVLGGPPVCSDATLPQ
		L				IINALEEDPAAQKMQLAF
16174	46542	Α	16269	120	327	
16175	46543	A	16270	540	791	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion) DRVSLLPRLRQENRLNPGGRGC
						SELRSHHCTPAWATE*GSVSKK KKKKKNQKKNQFDFISLALPPS PHCPPLSQSLLLQPSSSAN
16177	46545	A	16272	3	343	DIHHVRKSMEVNFLSYVVLTV AALPMLKQSNOSIVVVSSLAET AMKAVSGIVHMQAAPKEECAL EIIKGGALRQEEVYYDSSLWTT L/LIRNPCRKILEFLYSTSYNMD RFINK
16178	46546	В	16273	48	407	
16179	46547	А	16274	35	646	DLTGFIKFSLLIGLFMAYYYYS ANEEFRPGKYPCVSFWRNFFK KHRGAYAYHLAKMGAHVVV TARSKETLQKVRVLCSQICVPS HAQMCSYIPDMTAEQFVAQA GKLMGEA VSLTSSSELCPWGH QELLDIHHVRKSMEVNFLSYVV LTVAALPMLKQSNOSIVVVSSL AGK WDRDIGG*KKKNARDD AVQTPGSKSAI
16180	46548	А	16275	221	393	LLYPANCVYMRCPQMLHFQGI QRPILL/HVSSGGAIHLQTGVET EEMRSAIAPDPIPLT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
16181	46549	A	16276	2025	4146	SDEWIISAGSVSGDYNSSAAVS KNDILLGIEHISVESETNIHSSEE. FIKTAKSQLVEIEPATQNPKTSIS SYERLQALQETCKENKGVPKQ GDTFLSFSAACETDVSSVYPPEK FEFETSATGSSMQSGSELLLKE REILTAGKQPSDSBFSASLLAG GKSVAKTGPSSMQSGSELLLKE AYAQTQSSLFYSPSSPMSSDDE SEIEDEDLKVELQRLYREHIQEV VILQTTQNKELQELYERLRSHIQEV VILQTTQNKELQELYERLRSHIQEV SKYAKTGELDELYERLRSHIQEV KILRSRPQSLTHVDNGIVVIDPL VESNAASCQQSPASKKGMFTD DLHKLYDDWTKEAVGNSLIK SLNQLKQSQHLARSHVDHUVDELV SNAALSQQSPASKKGMFTD DLHKLYDDWTKEAVGNSLIK SLNQLKQSQHLKETENWNKVI QLESVYHREDEATWLPQLRAQ GFQEFLEKRCFGAVVPVDLLKI ARTVSRQQQQHLHRQLPFT AALGAYSGGAGPIRHHPAHH FHHHQGAQGWAMLGYSAI CQACAYPGGDGLELVWFPGC LGKDRLAEWGRECQAERCTAG LUVGHIVGNKGTPPRAWFLPPA CLGKCRLAEWGRECQAERCTAG LUVGHIVGNKGTPPRAWFLPPA CLSHCWRLALIRFRFFVYFQLEE
						SEYALYNIPSYYSSHALKLSWN GKNLTTNQFLMQEVAKQLGLI
16182	46550	A	16277		542	MRAPPKSGQLQHCRPSRGALE; GDLPWEINPLSSCSLLHEKDPP MTSGPQTNQPKEHLTNFKSGV RP/LQGRLPWSFTLSGKSRFSGE GASTPTPYIS/GAPIPYFRTPTSY LCAPIPYVRTPTSYLCALTPFPL FWRHIRTSKRLN/LQQPGIPPEP, PPG/CLLQVPEI*PPGGQMPAAQ
16183	46551	A	16278	3	351	VRPRRDACLGPSPLAASPAFLG KGQVPQPLISLCPDPLFPHPNLE LRPNPLCPHPDL/GISVP*PLSRF SGGT*ELPNA*TAAA/AAF/LPE PPPG/CLLQVPEI*PPGQGMPAA QDSS
16184 16185	46552 46553	A	16279 16280	774	513 1007	ARSVTSSSELCGSWCLQASRHT HVPWCQL*KLPVVHLVQPQPC RGPAPGAALPAGSLEMPGTTEF ERGCHSPGLGIS

SEO ID	ICEO ID NO:	Mat	SEQ ID NO:	Nucleotido	Nucleatide teention of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ļ ·			sequence	' '	,
16186	46554	A	16281	507	520	LLLGLAAED*CCLIASEAP*TQS
10180	46334	^	10201	307	320	LFGGLFTRTHMKFGAVTQIRGP
1						PLGDKSPVLLLFALERORRHVL
1			l			
i						SMDPKLRCWSRTGKAAFPWCL
						IIAEMPDYSPTFQRCQTTQGRLP
		1				WSFTLSSKSRFSGEGARACYKC
		Į				QKSDHQAKECPQPGIPPKPCPIC
Ì		ŀ				AGPHWKSDCSTHLAATPRAPG
	ļ	Ļ				TLAQGSLTDSFLA
16187	46555	Α	16282	269	607	RARSEGAGLWSVVAPSAVSVFF
	1	1				VSDPRCAPFHRSPSCCSPVDRV
ļ.	1	ł				CRCLRRVALAPAALVLVCSGSH
ŀ	1	l				SQSPWNSGLKLSDRLLLGLAAE
						D*RCPIASEAP*TTTDAELTIPGL
16188	46556	С	16283	I	1752	
16189	46557	В	16284	146	372	
16190	46558	Α	16285	336	753	ARTPLAAASRGSGGTRGGAAW
						VAGDDLRGPAPGPAEPGPARRR
						RGKGTPSSRPRTQRRPRSAAMH
						GHRCPQPARRSREHPNTYPAGR
			ļ			RPPELRRGRSGVPPPRP*RRPGE
i		l			1	APRAPRPRAAEL*ARLPRPSVRP
						APARRV
16191	46559	Α	16286	1	768	FDELREEGFRQSVITNFSELKKD
1						V*THRKEAKNLEKK\LDEWLTR
		1	ĺ			INSVEETLNDPMELKTMAQELR
ł		1				DACTT*FREKRVKRNEQSLQEI
		1				WDYVKRPNLRLIGVPESDGEN
						GTKLENTLLDMIQENFPKLARQ
1						ANIQIQEIQQIPQRYSSRRATPR
1		1				HIIVRFTKVEMKEKMLRAAREK
		l	ŀ	1		GWVTHKGKPIRLTADLSAETLQ
						ARREWGPIFNILKEKNFQPRISY
		1				PAKLSFISEGEIKYLTDKQMLR
		1				DFVTTRPALKEL
16192	46560	Α	16287	345	894	EQRHNIPESLGHI*STSAGDTPA
						DRVWSGPPANSSRPAAEGPDC*
		1			i	KEN*QRARTS\NQNPICTSPSSK
1						TKEAKNLEKRLDEWLTRINSIE
		1				NTLKDLMELKTRARELRDACR
1	1	1		I	1	SFSSQFDQVEERVSVIEDQMNE
1	1	1			1	MKKEEKFREKRVKRNEQSLQEI
	1	1				WDSVKRPNLHLIGVPESDGENG
1	1	1		I	1	IKLENTL
16193	46561	A	16288	1	1206	
1.0175	1.0001	1		ı <u>. </u>		

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	l	
16194	46562	A	16289		882	MAPVSASEDFWQCPETSLISQL
1015	10302	ľ.	10207	ľ	1002	ESAPGIOLMEARDADKSLIMHE
		1				IAPTSNRELPVHKISIAKPCSML
		1				MISWFFVNTGOSFORKDOAAIF
	l					AVLOPLLVIHROTGSGVDLOOT
						PTDLOLRDLTVRRKNNKOKGI
	l					ASTSRKRTSTPKPHLODHSSSPA
						MEOSWMOKDFDKLAEVGFRR
	ł					SVITNFSELKEDVOTHHKEAKN
						LEKRLDEWLTRINSVENTLNDL
1						MELKTMAQELHDACTSFSRQF
						DOVEERVSVIEDKISEIKQEEKF
		1		İ		REKRVK\RNEOSLOTIWDYVKR
		l				PNLHLIGVS
16195	46563	A	16290	h	1008	MVRFGDELGGRYGGPGGERA
10193	40303	A	10290	l'	1008	RGGGAGGAGGPGPGGLQPGQR
		l				VLYKQSIAQRARTMALYNPIPV
		l				KONCFTVNRSLFVFSEDNVVRK
ŀ	į.	l				YAKRITEWPPFEYMILATIIANC
	1					IVLALEOHLPDGDKTPMSERLD
		1				DTEPYFIGIFCFEAGIKIIALGFV
1	1	1		1		FHKGSYLRNGWNVMDFVVVL
	1					TGRKAGLGCCGSGVESGGWGD
	1]	RSSSPAMEOSRMENDFDELTEV
		i				GFRKSVITNFSELKEDVRTHRK
		1			1	EAKNLEKRLDEWLTRMNSVEK
		1				TLNDLMELKTMARELRDACTS
		1		ł		FSSOFDQVQEMVSVIQDQISEM
İ				l		KREEKFREKRVKRNE/OSLO*I
		1				WDYVERPNLRL
16196	46564	В	16291	1	1257	WBT VEIG NEIGE
16197	46565	A	16292	3	467	KTPFVGYOHORPKVDKTTKMG
1017/	70303	1	1.02,2	ľ	1	RNOSRKAENSKNOSTSSPPKDP
	1		1			GSLPATEQSWTENDFDELTEVG
	1					FRRSAITNFSELKEHVLTRRKEA
		1				KNLEKRLDEWLTRINSVEKTLN
						DLMELKTMAOELRDARTSFNS
						*FNOVEKRISVIEDOIDEIK*EDK
16198	46566	c	16293	115	498	THE TERRISTIED GIDEIN EDR
10128	40300	<u>_</u> ب	10293	1113	1770	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16199	46567	Α	16294	2	1455	TERDSININKKDIYTKTQSVGRQ
						HQRPKVDKTTKMGRNQNRKA
	į.					ENSKNQSASSPPKDCSSSPAME
	ì					OSWMENNFDELTEIGFRRSVIT
						NFSELKEDVQTHCKEAKNLEK
						K\LDEWLTRINSIEKNLNDLMEI
		1			ļ.	KTMAQELRDACTSFSSQFDQVE
		l			l	ERVSVIEDOMNEMKQEEKFRE
						K\RVKRNEOSLOEIWDCVKRPN
						LHLIGIPESDGENGTKLDNTLQ
						DII*ENFPHLAR*ANIQIQEIQEIQ
			i			RTPORYSPRRATPRHIIVRFTKV
		1				DMKEKMLRAVREKGFKPTKIR
1		1				RDKEGHYIMVKGSIQQEELTIL
	l					NIYAPNTGGPRFIKQVLSDLQR
						DLDSHTIIMGDFNTPLSTLDRS
		l				MRQKINKDIQELNSALHQVDLI
l						DIYRTLOPKSTEYTFFSAPHRTY
		1	8			SKIDHVVGSKTLLSKCKRPETT
						TNCLSNRSAIKLELRIKKLTONR
1		1				SAAWKLNNLLLNDYWVHNEM
		1				KAEIKMFFETRTKIOHARISGIL
16200	46568	A	16295	395	656	FOLORPASCSSPIOHRKHIPRKP
10200	40,000	l^	10293	3,5	050	A\SQSSLPHTGKSARQPQEPQNR
						ENCLLRILIRVASEATRRWTRPS
						EFSSRSENPRAHHPRRALR
I6201	46569	A	16296	671	1126	RRKVSELEHRRPVWKCPRRPPE
10201	40309	l^.	10230	1071	1120	VRKPESARVSFSPRGGARRSRP
						GATRSPQPRVPAAASASFGFGG
						LGRGGRGERARRGRGPKLEGP
		l		1		ESSAVLA*RNFPSRPRRRRPGRA
		1				GVEIARLGGTGSSPRPCPKPDPT
		1				ASLSELFMIPSDIPRRRN
16202	46570	A	16297	2571	2716	THGMLEKRCCLOENMPIIPGKD
10202	40370	l^	10257	2371	2710	FMTKTLKAIATKVKIGK*DLIKS
			l			FCT PERMITTENED PERMIT
16203	46571	В	16298	54	576	TC1
16203	46572	A	16299	475	719	VTWIRNNNACFTLNWFYHERS
10204	40372	l^	10233	1775	/'''	HIWILKSTLOCGOIIIGNDVESR
1		1				HK\CPKPS*LVGSVELEIADNVL
l		1				POKFPAAKTIFALYF
16205	46573	A	16300	395	830	LSPCHNSFFFFFQAFDDIAKYFS
10203	40373	^	10300	373	0.50	KKEWEKMK*ATVPGPSLPLKK
		1		1	1	*TFCSFLGFKVTLPPFMRSVSLE
		1				FGNLYQQRKILM*L*STVEMSLI
		1			1	LFICDLHIVKIMPKKPAEEENGL
		1				
						KEVPEASGPQNDGKQLCPPGNP
	10001	<u> </u>	1.0001	54	504	STLEKINKTS
16206	46574	В	16301	54	576	

SEQ ID	SEQ ID NO:	Met				Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
16207	46575	A	16302	475	719	VTWIRNNNACFTLNWFYHERS
						HIWILKSTLOCGOIIIGNDVESR
						HK\CPKPS*LVGSVELEIADNVL
					ŀ	PQKFPAAKTIFALYF
16208	46576	A	16303	393	822	LSPCHNSFFFFFQAFDDIAKYFS
		1			1	KKEWEKMKSATVPGPSLPLKK
				ļ.		*TFCSFLGFKVTLPPFMRSNSLE
		1				FGNLYQQRKILMYF*STVEMSL
						ILFICDLHIVKIMPKKPAEEENG
						LKEVPEASGPONDGKOLCPPGN
						PSTLEKINK
16209	46577	A	16304	1	447	
16210	46578	A	16305	75	565	LLFSGFSLLHGYLPHWYLCHLP
10210	1.007.0	1	1	-		RSRSLGLRDKRWKVKGDTDFL
	!					SPSHPEYLLKEENDGHLCSLSFR
	1					MGNOFSSPLPAYTPLECILNHW
			l	i		DGFYP*NLEEKHLIALRTKGWP
	l .	i				NYDLOEGLSWPOEGTIHFNTT
						WOLELFCRHEDRWSEATYMO
		1				AFYILQGNRDLG
16211	46579	Α	16306	1	1620	
16212	46580	Α	16307	128	223	
16213	46581	Α	16308	1	1578	
16214	46582	В	16309	592	883	
16215	46583	Α	16310	1	344	MHQTTAKTPHTSYPNNYEEHN
		ļ				RRANRSKNYDRIRHTTIILKRTN
		l	l			PAQTITNLLKSRFNRYNYVQPR
		1				SCRSIFIHIPGERDLNLCIAPNPP
		1	l			AIFPTSIGYLGPDLNNLPTYAPFI
		1	l	1		TTNLTGPAPTTTHPVGNLGTTG
	1					NPEPGGCGILIVWCCVNWACL
	l	[WDSVVAAPREPYMPAGLLRHR
l	Ì		ŀ			RHYRHRREDLIAVPAARQTDE
	ł	1				NRRPALCHEYAVGHRKIKTNV
l		ı				EHTEPYILPLITHSQPLNRGAFF
		1				CPGVRFLVVSVLDFEKRLAFAS
		ı	i			*SCGVOLRIRGKVOGVGFRPFV
		1	l		1	WOLAOOLNLHGDVCNDGDGV
		1				EVRLAYRVLVGOPOHYPRGRP
	I	A	16311	84	237	DGIVLSTYYHFYWHHF*ILEGP
16216	146584					
16216	46584	l ^A	10311	84	231	LPANTGDLCLPLOVOPDREOVR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
16217	46585	Α	16312	2887	3537	HLDRSGLFNIQVIPVCRKLGKE
						GNDEKQCVTTSRSNAAFFLLHF
						FLLATFTWMGLEAIHMYIALVK
						VFNTYIRRYILKFCIIGWGKPLK
						IFLVLFFPHENCOVYGKESYGK
				i		EKGDEL*VIKTFCDE*ISFVS*SG
ŀ				İ		VMFFLNIAMFIVVMVQICGRNG
						KRSNRTLREEVLRNLRSVVSLT
-		i		l		FLLGMTWGFAFFAWGPLNIPF
						MYLFSIFNSLQGKINCT
16218	46586	Α	16313	302	547	QIGLWTPNFRARTFCVWTGGL
				I		VLCF*GFCFQEAAGARDTSSFA
						MGVSPWEFRHG*LWPEMRSME
				ł	1	QILELLVLEEFLTITGQ
16219	46587	Α	16314	1	1174	WEEKENKEYLKSLFEILILMGK
						QNIPLDGHEADEIPEGLFTPDNF
						QALLECRINSGEEVLRKRFETT
						AVNTLFCSKTQQRQMLEICESC
						IREETLREVRDSHFFSIITDDVV
İ						DIAGEEHLPVLVRFVDESHNLR
						EEFIGFLPYEADAEILAVKFHTM
						ITEKWGLNMEYCRGQAYIVSS
						GFSSKMKVVASRLLEKYPQAIY
						TLCSSCALNMWLAKSVPVMGV
						SVALGTIEEVCSFFHRSPQLLLE
	ł					LDNVISVLFQNSKERGKELKEIC
	ì					HSQWTGRHDAFEILVELLQALV
					1	LCLDGINSDTNIRWNNYIAGRA
						FVL\CSAVSDFDFIVTIVVLKNV
				i		LSFTRAFGKNLQGQTSDVFFAA
	l					GSLTAVLHSLNEVMENIEVYHE
						FWFEEATNLAHQT
16220	46588	Α	16315	717	902	GALQSWPCYLPRQCGATVQNF
						CGVPTCMRKSTWSNLVVFRFP
		_				QDPSRCAN**LDCCTACTQ
16221	46589	Α	16316	1	374	EHSEPPIGVQIVDYLLRQEKVT
		ł				DRMDHSKVETETVLSFVDDIIS
						GAKSPCAMPSQVPDK/QAQEIA
l	l	1		l		DKIYNLFNGYTSGKEQQTAYN
		ĺ				TLLDLGSPTLHRVLYHYNQHY
L						ESFGEFTWRCEDELGPR
16222	46590	Α	16317	2	1293	